ca Hi 45	S GI	g to n Ti	39 ga Tp G]	ia aa .u As	t ac n Th 46	r Pro	t tta	a aa u Ly	a gc s Al	a gaa a Glu 469	u Ar	a ca g Hi	it ag .s Ai	ja a :g Tl	et gaa nr Glu 470	·
ga As 47	ь гл	g aç s Ar	g Ly	a ag s Ar	a ga g Gl: 47	u Glr	a gaa n Glu	a ag	c aaa r Lys	a gaa s Glu 481	ı Gl	a aa u As	t ag n Ar	g ca	at att is Ile 486	:
ag Ar 48	g As	t ga n Gl	a aa u Ly	a ag s Ar	a gta g Val 492	l Pro	aca Thr	a gaa Glu	a cat u His	t ttg Leu 497	ı Glı	g aa n Ly	g ac s Th	t aa r As	t aag n Lys 502	1720
503	ı ını	с гу	в гу	s Th	508	Thr	Asp) Let	ı Lys	513	Glr	í Ası	n Gl	u Pr	a aag o Lys 518	1768
519	ASI	о гу	s GT	y GI	1 Val 524	. Leu	Asp	Asn	ı Gly	7 Val 529	Ser	Glı	ı Gl	y Al	a gat a Asp 534	1816
535	гг	G G-11	ı Lei	ı Ala	Met 540	Lys	Ala	Glu	Ser	Gly 545	Pro	Asr	ı Glı	ı Th	a aaa r Lys 550	1864
551	пys	Asp) Let	ı Lys	556	Ser	Phe	Met	Lys	Lys 561	Leu	Asn	Let	ı Th:	t ctt Leu 566	1912
567	Pro	Ala	. Буз	Lys	Gln 572	Pro	Val	Ser	Gln	Asp 577	Asn	Gln	His	Lys	ata Ile 582	1960
583	Asp	тте	Pro	гуs	588	Ser	Gly	Val	Cys	Asp 593	Ser	Glu	Ser	Ser	atg Met 598	2008
599	vai	ьys	Thr	Val	A1a 604	Tyr	Val	Pro	Ser	Ile 609	Ser	Glu	His	Ile	ttg Leu 614	2056
615	GIU	Ala	Ата	Val	agt Ser 620	Glu	His	Thr	Met	Gly 625	Glu	Thr	Lys	Ser	Thr 630	2104
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647	116	ser	GIU	Thr	aac Asn 652	Lys (Glu /	Asp	Glu .	Asn 9 657	Ser	Leu	Leu	Val	Arg 662	2200
663	vai	Asp	Asn	Tnr	atg Met 668	His (Cys (Glu (Glu 1	Pro 1 673	[le	Cys	Gly	Thr	Glu 678	2248
act	CCC	CTC	cca	tct	cct .	atg g	gaa a	ata (caa (cag a	ıca 🤉	gaa	tcc	ttg	ttt	2296

Thr 679	Ser	Phe	Pro	Ser	684	o Mei	Gl	u Il	e Gl:	n Gl 68		r Gl	u Se	r Le	u Phe 694	
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cct Pro 711	gtg Val	gta Val	atg Met	gat Asp	gta Val 716	. Let	caa Gli	a aca	a gat c Asp	gte Va: 72:	l Se	t caa r Gli	a aa n As:	c tt n Ph	t ggc e Gly 726	2392
ttg Leu 727	gaa Glu :	ttg Leu	gat Asp	acc Thr	aaa Lys 732	Arg	aat Asr	gat 1 Asp	aat Asr	tca Ser 737	: As	t tat o Tyr	tgi Cys	gg Gl	t att y Ile 742	2440
tct Ser 743	gaa g Glu (ggt 3ly	atg Met	gag Glu	atg Met 748	Lys	gto	gca Ala	ctt Leu	tca Ser 753	Th	a aca r Thr	gtg Val	g agi	gaa Glu 758	2488
acc a Thr 7	act o	gaa Slu	agc Ser	att Ile	ttg Leu 764	Gln	cct	tca Ser	att	gag Glu 769	Gli	a gct 1 Ala	gat Asp	att Ile	ttg Leu 774	2536
cca a Pro] 775	ata a Ile M	atg (Met 1	ctt Leu	tca Ser	gaa Glu 780	gat Asp	aat Asn	aac Asn	cca Pro	aaa Lys 785	Phe	gag Glu	cct	tct Ser	gtt Val 790	2584
ata g Ile V 791	gtt a /al T	ca d hr I	cca Pro	ctg Leu	gtt Val 796	gag Glu	agt Ser	aag Lys	tcg Ser	tgt Cys 801	cat His	ctg Leu	gag Glu	cct	tgc Cys 806	2632
tta c Leu P 807	ct a Pro L	aa c ys C	gag : Blu '	act Thr	cta Leu 812	gat Asp	tct Ser	tca Ser	ctt Leu	cag Gln 817	cag Gln	act Thr	gag Glu	tta Leu	atg Met 822	2680
gac c Asp H 823	ac a is A	ga a rg M	itg g let <i>l</i>	Ala	act Thr 828	ggt Gly	gaa Glu	aca Thr	aac Asn	tca Ser 833	gta Val	tat Tyr	cat His	gat Asp	gat Asp 838	2728
gat a Asp A 839	ac te sn Se	cg g er V	tt t al I	Leu	agc Ser 844	att Ile	gac Asp	ctt Leu	aat Asn	cac His 849	ctg Leu	aga Arg	cct Pro	att Ile	cca Pro 854	2776
gaa g Glu A 855	cc at la I	tc a le S	gt o er F	ro :	ctg Leu 860	aat Asn	agt Ser	cca Pro	Val	aga Arg 865	cct Pro	gta Val	gca Ala	aaa Lys	gtt Val 870	2824
ctt ag Leu A 871	ga aa rg As	at g sn G	aa a lu S	er :	cca Pro 876	cct Pro	caa Gln	gtt Val	Pro	gtg Val 881	tat Tyr	aat Asn	aac Asn	agt Ser	cat His 886	2872
aaa ga Lys As 887	at gt sp Va	g t	tt t he L	eu I	cca Pro 1	aat Asn	tca Ser	gct Ala	His	tct Ser 897	acc Thr	tct Ser	aag Lys	agt Ser	cag Gln 902	2920
tct ga Ser As	at ct sp Le	c aa u As	at a sn L	ag g ys C	gaa a Blu A	aat (Asn (caa Gln	aag Lys :	cca Pro :	att Ile	tac Tyr	aaa Lys	tct Ser	gac Asp	aaa Lys	2968

	903	;				908	3				913	3				918	
	tgt Cys 919	Thr	gaa Glu	gca Ala	gac Asp	aca Thr 924	Cys	aag Lys	g aat s Asn	tca Ser	cca Pro	Leu	gat Asp	gaa Glu	tta Lev	gaa Glu 934	3016
	gaa Glu 935	Gly	gaa Glu	att	aga Arg	agt Ser 940	Asp	agt Ser	gaa Glu	aca Thr	tct Ser 945	Lys	. cca Pro	caa Gln	gaa Glu	agt Ser 950	3064
	ttt Phe 951	Glu	aaa Lys	aat Asn	tcc Ser	aaa Lys 956	Arg	aga Arg	gtg Val	tca Ser	gct Ala 961	Asp	gtg Val	cgg	aag Lys	tca Ser 966	3112
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terif-stire then thin that tery	agt Ser 983	agg Arg	aaa Lys	aca Thr	cat His	gta Val 988	aga Arg	atc Ile	cat His	cag Gln	acc Thr 993	aat Asn	aac Asn	aaa Lys	tgg Trp	aat Asn 998	3208
Harff-Stires Assess His	aaa Lys 999	aga Arg	cct Pro	gat Asp	aaa Lys	tct Ser 1004	agt Ser	aga Arg	tct Ser	Ser	aaa Lys 10.09	aca Thr	gag Glu	aag Lys	Lys	gat Asp 1014	3256
ilmi ilmi	aaa Lys 1015	gtg Val	atg Met	agc Ser	act Thr	tcc Ser 1020	agc Ser	ttg Leu	gaa Glu	Lys	ata Ile 1025	gtt Val	cca Pro	att Ile	Ile	gct Ala 1030	3304
freer tente if it thereft	gta Val 1031	ccc Pro	tct Ser	tct Ser	gaa Glu 1	caa Gln 1036	gag Glu	atc Ile	atg Met	His	atg Met 1041	tta Leu	cga Arg	atg Met	Ile	aga Arg 1046	3352
	aaa Lys 1047	cat His	gta Val	aga Arg	aaa Lys 1	aat Asn .052	tat Tyr	atg Met	aaa Lys	Phe	aag Lys 1057	gca Ala	aaa Lys	ttt Phe	Ser	tta Leu 1062	3400
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	ttt Phe 1143	: Ala	aa <u>c</u> Lys	ctt Lev	Lys	aaa Lys 1148	Ile	tta Leu	gta Val	tgt Cys	gat Asp 1153	Ser	aaa Lys	ago Sei	ttt Phe	gga Gly 1158		3688
		Asp			Glu		Lys					Ser				gca Ala 1174		3736
	cag Gln 1175	Tyr	tca Ser	aat Asn	agt Ser	cag Gln 1180	Lys	agg Arg	agt Ser	gtg Val	gac Asp 1185	aac Asn	tcc Ser	aac Asn	aga Arg	gaa Glu 1190		3784
	ttg Leu 1191	Leu	aaa Lys	gaa Glu	aaa Lys	tta Leu 1196	Ser	aaa Lys	tca Ser	Glu	gac Asp 1201	cct	gtt Val	cat His	tat Tyr	aag Lys 1206		3832
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	tct Ser 1207	Leu	gtg Val	gga Gly	tgt Cys	aaa Lys 1212	aaa Lys	tct Ser	gag Glu	Glu	aat Asn 1217	tat Tyr	caa Gln	gac Asp	Gln	aat Asn 1222		3880
tions from Give Ame that 4m4	aac Asn 1223	tcc Ser	agt Ser	att Ile	aac Asn	act Thr 1228	gta Val	aag Lys	cat His	Asp	att Ile 1233	aaa Lys	aaa Lys	aat Asn	Phe	aac Asn 1238	-	3928
	atc Ile 1239	tgc Cys	ttt Phe	gat Asp	aat Asn	ata Ile L244	aag Lys	aac Asn	tct Ser	Gln	tcc Ser 1249	gaa Glu	gag Glu	cgc Arg	Ser	ttg Leu 1254	٠	3976
t that is it there is the	gaa Glu 1255	gta Val	cac His	tgt Cys	cca Pro 1	agc Ser 1260	acc Thr	cca Pro	aag Lys	Ser	gaa Glu 1265	aaa Lys	aac Asn	gaa Glu	Gly	agc Ser 1270		4024
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	gag Glu 1335	tgg Trp	gag Glu	tta Leu	aag Lys 1	act Thr 340	cca Pro	gag Glu	aag Lys	Gln	ctg Leu 345	cta Leu	gag Glu	act Thr	Leu	aag Lys 350		4264
	tgc Cys 1351	gag Glu	tct Ser	ata Ile	cca Pro 1	gct Ala 356	tgt Cys	aca Thr	aca Thr	Glu	gag Glu 361	cta Leu	gtt Val	tca Ser	Gly	gtg Val 366	•	4312

	gct Ala 1367	tct Ser	cca Pro	tgt Cys	Pro	aaa Lys 1372	Met	att Ile	agt Ser	Asp	gat Asp 1377	Asn	tgg Trp	tca Ser	tta Leu	tta Leu 1382	4360
					Gly		Ser			Ser						gtt Val 1398	4408
					Leu		Glu			Met						aac Asn 1414	4456
					Lys		Asn			Ser						ccc Pro 1430	4504
of the					Ile		ctt Leu			Leu					Thr		4552
men them down thank					Lys		gat Asp			Leu					Pro		4600
land that then					Ser		cct Pro			Val					Phe		4648
then? B it graft 19° that					Leu		gaa Glu			Ala					Ile		4696
them then?					Ser		aat Asn			Ser					Ser		4744
					Arg		gtt Val			Gly					Pro		4792
					Ala		ata Ile			Lys					Phe		4840
	gtg Val 1543				Arg		aca Thr			Thr					Lys		4888
		_	_		Asp	_	tta Leu	_		Ser	_		_		Gly	_	4936
					Gly		gag Glu			Tyr					Asn		4984
	gtt	ttt	aaa	tct	gtg	gag	gaa	ttg	gaa	aac	tcc	aac	aaa	aat	gtt	gat	5032

Val Phe	Lys Ser Val G	lu Glu Leu Glu	Asn Ser Asn Lys	Asn Val Asp
1591	15	96	1601	1606
ggc agc	aag tca act c	is Giu Giu Gir	g agc tct atg ata	caa aca cag 5080
Gly Ser	Lys Ser Thr H		Ser Ser Met Ile	Gln Thr Gln
1607	16		1617	1622
gtt cct	gat ata tat ga	lu Phe Leu Lys	gat gct tca gat	aag atg ggt 5128
Val Pro	Asp Ile Tyr G		Asp Ala Ser Asp	Lys Met Gly
1623	162		1633	1638
cat agt	gat gaa gtg go	a Asp Glu Cys	ttc aaa ttg cat	caa gta tgg 5176
His Ser	Asp Glu Val Al		Phe Lys Leu His	Gln Val Trp
1639	164		1649	1654
gaa aca a	aaa gtg cct ga	u Ser Ile Glu	gaa ttg cct tca	atg gaa gaa 5224
Glu Thr 1	Lys Val Pro Gl		Glu Leu Pro Ser	Met Glu Glu
1655	166		1665	1670
atc tca c	cac tct gtt gg	y Glu His Leu	cca aac aca tac	gta gat cta 5272
Ile Ser I	His Ser Val Gl		Pro Asn Thr Tyr	Val Asp Leu
1671	167		1681	1686
acg aaa g	gat cca gtc ac	r Glu Thr Lys	aac ttg ggg gaa	ttc ata gaa 5320
Thr Lys A	Asp Pro Val Th		Asn Leu Gly Glu 1	Phe Ile Glu
1687	169		1697	1702
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Val Thr V	al Leu His Ilo		Gly Cys Ser Gly (Gly Asn Leu
1703	170		1713	1718
aat cag a	gt gct caa ata	Leu Asp Asn	tct ttg cag gct g	at act gta 5416
Asn Gln S	er Ala Gln Ile		Ser Leu Gln Ala A	sp Thr Val
1719	1724		1729	1734
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Gly Ala P	he Ile Asp Lei		Ala Ser Ser Glu A	la Lys Ser
1735	1740		1745	1750
gaa ggt aa	at cat cct gca	Leu Ala Val	gaa gac ttg gga t	gt ggg gtg 5512
Glu Gly Aa	sn His Pro Ala		Glu Asp Leu Gly C	ys Gly Val
1751	1756		1761	1766
ata cag gt	ta gat gaa gat	aat tgt aag o	gaa gaa aag gca c	aa gtg gca 5560
Ile Gln Va	al Asp Glu Asp		Glu Glu Lys Ala G	ln Val Ala
1767	1772		1777	1782
aac agg co Asn Arg Pr 1783	t tta aaa tgc O Leu Lys Cys 1788	att gtt gag g Ile Val Glu (gaa acc tat atc ga Flu Thr Tyr Ile As 1793	ac ttg acc 5608 sp Leu Thr 1798
aca gaa to Thr Glu Se 1799	et ccc agt tca er Pro Ser Ser 1804	tgt gaa gta a Cys Glu Val I	aaa aaa gat gag tt ys Lys Asp Glu Le 1809	a aaa tca 5656 u Lys Ser 1814
gag cca gg	a tca aat tgt	gat aac tcg g	ag ttg cct ggg ac	t ttg cat 5704
Glu Pro Gl	y Ser Asn Cys	Asp Asn Ser G	lu Leu Pro Gly Th	ur Leu His

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123 123
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1815					1820					1825	;				1830	
aat Asn 1831	tct Ser	cac His	aaa Lys	Lys	aga Arg 1836	Arg	aac Asn	att Ile	Ser	gat Asp 1841	Leu	aat Asr	cat His	cct Pro	cat His 1846	57
aaa Lys 1847	aaa Lys	caa Gln	aga Arg	Lys	gaa Glu 1852	aca Thr	gac Asp	tta Leu	Thr	aat Asn 1857	Lys	gaa Glu	aag Lys	acc Thr	aag Lys 1862	58
aaa Lys 1863	cct Pro	acc Thr	caa Gln	Asp	tct Ser 1868	tgt Cys	gag Glu	aat Asn	Thr	gaa Glu 1873	gct Ala	cac His	caa Gln	aag Lys	aaa Lys 1878	58
gcc Ala 1879	agt Ser	aag Lys	aag Lys	Lys	gcc Ala 1884	cct Pro	cct Pro	gtg Val	Thr	aaa Lys 1889	gat Asp	ccc Pro	tca Ser	tca Ser	tta Leu 1894	58
aag (Lys 1 1895	gca Ala	acc Thr	cca Pro	Gly	att Ile 1900	aag Lys	gat Asp	tca Ser	Ser	gca Ala 1905	gca Ala	ctt Leu	gcc Ala	Thr	tct Ser 1910	59
aca a Thr 9 1911	agt Ser	ctt Leu	tct Ser	Ala	aaa Lys 1916	aat Asn	gtt Val	att Ile	Lys	aag Lys 1921	aag Lys	gga Gly	gaa Glu	Ile	atc Ile 1926	59
att t Ile I 1927	tta Leu	tgg Trp	aca Thr	Arg	aat Asn 932	gat Asp	gac Asp	cgg Arg	Glu	att Ile .937	tta Leu	ttg Leu	gag Glu	Cys	cag Gln 1942	60-
aaa a Lys A 1943	aga Arg	gly 999	cca Pro	Ser	ttt Phe 948	aaa Lys	aca Thr	ttt Phe	Ala	tat Tyr 953	tta Leu	gcc Ala	gcc [.] Ala	Lys	ttg Leu L958	608
gat a Asp I 1959	aaa Lys	aat Asn	cca Pro	Asn	cag Gln 964	gtc Val	tca Ser	gaa Glu	Arg	ttc Phe 969	cag Gln	cag Gln	cta Leu	Met	aag Lys 1974	613
ctc t Leu P 1975	he	gaa Glu	aag Lys	Ser	aaa Lys 980	tgc Cys	agg Arg	tag *	ttaa	t gg	ttac	acta	caç	gaga	ıcta	618
gtaaa	ata	aa t	gttt	gaca	c tg	tagt	tgtg	cag	ttaa	tgg -	cctg	ttag	tc a	ttga	agatg	624
tgaat	agt	gg g	tgag	agac	a tc	cagt	tgag	ttc	ctgc	ttc	agtc	acag	gt t	ggag	gatga	630
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<212> DNA

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<220>

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Ser 154	T His	s Se	r Sei	c Glu	1 Glu 159	ı Asp	Glı	ı Ala	a Ser	Ser 164	Glr	n Thi	As)) Le	a agc u Ser 169	951
Caa Glr 170	Thi	g ato	c tca e Sei	a aag	g aaa E Lys 175	Thr	gto Val	agg L Arg	g ago g Ser	: ata : Ile 180	Glr	a gag n Glu	g gct 1 Ala	e cc	a gtg o Val 185	999
agt Ser 186	Glu	a gat a Asp	t ctt	gta Val	ato Ile 191	Arg	tta Lev	a cgt a Arg	cga Arg	Pro 196	Pro	cta Leu	aga Arg	tao Tyi	c cca Pro 201	1047
Arg 202	Tyr	Glu	ı Ala	Thr	Ser 207	Val	Gln	Gln	Lys	Val 212	Asn	Phe	Ser	Ğlı	gaa Glu 217	1095
gga Gly 218	Glu	act Thr	gaa Glu	gaa Glu	gat Asp 223	gat Asp	caa Gln	gac Asp	agc Ser	tct Ser 228	cac His	agc Ser	agt Ser	gto Val	act Thr 233	1143
Thr 234	Val	Lys	Ala	Arg	Ser 239	Arg	Asp	Ser	Asp	Glu 244	Ser	Gly	Asp	Lys	acc Thr 249	1191
Thr 250	Arg	Ser	Ser	Ser	Gln 255	Tyr	Ile	Glu	Ser	Phe 260	Trp	Gln	Ser	Ser	caa Gln 265	1239
Ser 266	Gln	Asn	Phe	Thr	Ala 271	His	Asp	Lys	Gln	Pro 276	Ser	Val	Leu	Ser	281	1287
Gly 282	Tyr	Gln	Lys	Thr	Pro 287	Gln	Glu	Trp	Ala	Pro 292	His	act Thr	Ala	Arg	Ile 297	1335
Arg 298	Thr	Arg	Met	Gln	Asn 303	Asp	Ser	Ile	Leu	308	Ser	gag Glu	Leu	Gly	Asn 313	1383
Gln 314	Ser	Pro	Ser	Thr	Ser 319	Ser	Arg	Gln	Val	Thr 324	Gly	caa Gln	Pro	Gln	Asn 329	1431
Ala 330	Ser	Phe	Val	Lys	Arg 335	Asn	Arg	Trp	Trp	Leu 340	Leu	cct Pro	Leu	Ile	Ala 345	1479
A1a 346	Leu	Ala	Ser	Gly	Ser 351	Phe	Trp	Phe	Phe	Ser 356	Thr	cct Pro	Glu	Val	Glu 361	1527
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Gly Glu Ser Ala Ser Asp Gly His Ile Ser Cys Pro Lys Pro Ser Ile
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tion them that their die	gac Asp 1064	Asn	aaa Lys	att Ile	cac His	att Ile 1069	cac His	tta Leu	gly ggg	Ser	cag Gln 1074	ttt Phe	aaa Lys	cgg Arg	tcc Ser	cct Pro 1079	3870
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Asn Pro Tyr Leu Ile Arg Arg Ile Glu Glu Leu Asn Lys Thr Ala Ser
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Series of the series	cag tgg tat Gln Trp Tyr 416	tct tgg gg Ser Trp Gl 42	y pro pro	Asn Met (cag tgt aga tta tgt Gln Cys Arg Leu Cys 126	gca 1536 Ala 431
	att tgt tgg (Ile Cys Trp) 432	ctt tat tg Leu Tyr Tr 43°	ь газ газ	Tyr Gly G	ggc ttg aaa atg ccc Gly Leu Lys Met Pro 142	acc 1584 Thr 447
and the state	cag tca gaa g Gln Ser Glu (448	gaa gag aag Glu Glu Lys 453	Leu Ser	Pro Ser P	ca act aca gag gac ro Thr Thr Glu Asp 58	cct 1632 Pro 463
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                                                                     2068
aaagaagctt aatataaaag tttgcattct actcaggaaa aagcatcttc ttgtatatgt
                                                                     2128
cttaaatgta tttttgtcct catatacaga aagttcttaa ttgattttac agtctgtaat
                                                                     2188
gcttgatgtt ttaaaataat aacattttta tatttttaa aagacaaact tcatattatc
                                                                     2248
ctgtgttctt tcctgactgg taatattgtg tgggatttca caggtaaaag tcagtaggat
                                                                     2308
ggaacatttt agtgtatttt tactccttaa agagctagaa tacatagttt tcaccttaaa
                                                                     2368
agaaggggga aaatcataaa tacaatgaat caactgacca ttacgtagta gacaatttct
                                                                     2428
gtaatgteee ettettteta ggetetgttg etgtgtgaat eeattagatt tacagtateg
                                                                     2488
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                                                                     2548
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Ala Lys Leu Glu Lys His Leu Met Leu Leu Arg Gln Glu Tyr Val Lys
                                                                      101
ctg caa aag aag ctg gcg gag aca gag aag cgc tgc gct ctc ttg gct
Leu Gln Lys Lys Leu Ala Glu Thr Glu Lys Arg Cys Ala Leu Leu Ala
                                                                      149
23
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gcg Ala 39	i Glr	g gca n Ala	a aad a Asr	c aag n Lys	g gag s Glu 44	ı Ser	ago Sei	c ago r Sei	gag Glu	g tco 1 Sen 49	r Ph	c ato e Ile	e Sei	c cg	t ctg g Leu 54	197
ctg Leu 55	ı Ala	ato a Ile	gto Val	g gca L Ala	a gad a Asp 60	Let	tac Tyi	gag Glu	g cag Glr	g gag n Glu 65	ı Glı	g tad	ago Sei	gat Asj	ctg Leu 70	245
aag Lys 71	Ile	a aag E Lys	gtt Val	ggg Gly	g gac Asp 76	Arg	cac His	ato Ile	agt Ser	gct Ala 81	His	c aag s Lys	ttt Phe	gto Val	c ctg Leu 86	293
tca Ser 87	Ala	cgc Arg	agt Ser	gac Asp	ago Ser 92	tgg Trp	agt Ser	ctg Leu	gct Ala	aac Asn 97	Let	tct Ser	tcc Ser	act Thr	aaa Lys 102	341
Glu 103	Leu	Asp	Leu	Ser	Asp 108	Ala	Asn	Pro	Glu	Val 113	Thr	Met	Thr	Met	ctt Leu 118	389
Arg 119	Trp	Ile	Tyr	Thr	124	Glu	Leu	Glu	Phe	Arg 129	Glu	Asp	Asp	Val	134	437
Leu 135	Thr	Glu	Leu	Met	Lys 140	Leu	Ala	Asn	Arg	Phe 145	Gln	cta Leu	Gln	Leu	Leu 150	485
Arg 151	Glu	Arg	Cys	Glu	Lys 156	Gly	Val	Met	Ser	Leu 161	Val	aat Asn	Val	Arg	Asn 166	533
Cys 167	Ile	Arg	Phe	Tyr	Gln 172	Thr	Ala	Glu	Glu	Leu 177	Asn	gcc Ala	Ser	Thr	Leu 182	581
Met 183	Asn	Tyr	Cys	Ala	Glu 188	Ile	Ile	Ala	Ser	His 193	Trp	gac Asp	Asp	Leu	Arg 198	629
Lys 199	Glu	Asp	Phe	Ser	Ser 204	Met	Ser	Pro	Gln	Leu 209	Leu	tac Tyr	Lys	Met	Ile 214	677
Lys 215	Ser	Lys	Thr	Glu	Tyr 220	Pro	Leu	His	Lys	Ala 225	Ile	aaa Lys	Val	Glu	Arg 230	725
G1u 231	Asp	Val	Val 	Ser	Leu 236	Tyr	Leu	Ile	Glu	Met 241	Asp	tcc Ser	Gln	Leu	Pro 246	773
999 Gly 247	aag Lys	ctg Leu	aạt Asn	gaa Glu	gcg Ala 252	gac Asp	cat His	aac Asn	Gly .	gac Asp 257	ctg Leu	gca Ala	tta Leu	Asp	cta Leu 262	821

gc: Al: 26:	a Lei	c tca 1 Sei	a cga r Arg	a cga g Arg	t cto Lev 268	Glu	g aat i Asr	att 1 Ile	gco Ala	a Caco a Thi 273	Thi	g cto	g gt	t ag l Se	t cac r His 278	869
aa: Ly: 27:	s Ala	gat Asp	gtg Val	g gad Asp	atg Met 284	. Val	gac Asp	aag Lys	g agt Ser	ggc Gl _y 289	Trp.	g ago Sei	tte	g tt u Le	a cac u His 294	917
aaa Lys 295	s Gly	ato / Ile	c caa e Gln	aga Arg	gga Gly 300	Asp	Ctc Leu	ttt Phe	gct Ala	gcc Ala 305	Thr	tto Phe	cto Lei	at Ille	t aag e Lys 310	965
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Arg 487		ı Ly	s Trj	p Gly	y Glu 492		Pro	o Lei	ı His	497		a Cys	s Arg	g Hi	s Gly 502	
cto Lev 503	ı Ala	c aa a As:	c cto n Lev	c aca	a gca Ala 508	ı Glu	cto Le	c ctg ı Lev	g cag ı Glr	g caa n Glr 513	ı Gl	c gcd y Ala	c aad a Asi	c cca	a aac o Asn 518	1589
cto Leu 519	Glr	g acg	g gag r Glu	g gaa 1 Glu	gct Ala 524	ı Leu	Pro	ctg Leu	cca Pro	aag Lys 529	Gli	g gco	gca Ala	a tco a Sei	c ctg Leu 534	1637
acc Thr 535	Ser	ttg Lei	g gcg ı Ala	g gac Nap	ago Ser 540	· Val	cat	ctg Leu	cag Gln	acg Thr 545	Pro	a ctg D. Leu V	cac His	ato Met	g gcg : Ala 550	1685
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gcc Ala 567	Asn	gct Ala	ctt Leu	cat His	gcc Ala 572	acc Thr	aac Asn	aac Asn	ttg Leu	cag Gln 577	Ile	: att : Ile	ccg Pro	gac	ttc Phe 582	1781
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gat Asp 647	ata Ile	aat Asn	gtc Val	agc Ser	agg Arg 652	act Thr	cag Gln	gac Asp	gly ggg	gag Glu 657	aca Thr	gcc Ala	ctc Leu	cag Gln	ctg Leu 662	2021
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ctt Leu 695	gca Ala	ttg Leu	gca Ala	aac Asn	aat Asn 700	ctg Leu	gag Glu	gac Asp	atc Ile	gca Ala 705	tcc Ser	act Thr	ctg Leu	gtc Val	aga Arg 710	²¹⁶⁵
cat His	ggc Gly	tgt Cys	gat Asp	gcc Ala	aca Thr	tgc Cys	tgg Trp	ggt Gly	ccg Pro	gga Gly	cct Pro	ggt Gly	999 Gly	tgc Cys	ctt Leu	2213

	711				716				721				726	
										Asn			gcc Ala 742	2261
		ttt Phe											cca Pro 758	2309
		gcc Ala												2357
a Ł		cat His												2405
Town the third	_	gag Glu		_			_	_	_	_	-	 _		2453
40 if all reads all than the tard that the		atc Ile		_		-				-		_		2501
		gtt Val												2549
Sand to the stands of the stands		acc Thr												2597
		gcc Ala												2645
		ggc Gly												2693
		gtg Val												2741
		gat Asp												2789
		gaa Glu												2837
		tta Leu												2885

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		Ala										His			gtc Val		2981
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	gac Asp 999	gcc Ala	gaa Glu	gcc Ala	Phe	aat Asn 1004	ctc Leu	aga Arg	ggc Gly	Gln	tca Ser 1009	cca Pro	ctg Leu	cac His	att Ile	ttg Leu 1014	3077
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	gaa Glu 1031	tgc Cys	atg Met	ccg Pro	Gly	tat Tyr 1036	cct Pro	ctg Leu	gac Asp	Lys	ccg Pro 1041	gat Asp	gca Ala	gac Asp	ggc Gly	agc Ser 1046	3173
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ctg caa aag aag ctg gcg gag aca gag aag cgc tgc gct ctc Leu Gln Lys Lys Leu Ala Glu Thr Glu Lys Arg Cys Ala Leu 23 28 33	ttg gct 149 Leu Ala 38
gcg cag gca aac aag gag agc agc agc gag tcc ttc atc agc Ala Gln Ala Asn Lys Glu Ser Ser Ser Glu Ser Phe Ile Ser 39 44 49	cgt ctg 197 Arg Leu 54
ctg gcc atc gtg gca gac ctc tac gag cag gag cag tac agc Leu Ala Ile Val Ala Asp Leu Tyr Glu Gln Glu Gln Tyr Ser 55 60 65	gat ctg 245 Asp Leu 70
aag ata aag gtt ggg gac agg cac atc agt gct cac aag ttt Lys Ile Lys Val Gly Asp Arg His Ile Ser Ala His Lys Phe 71 76 81	gtc ctg 293 Val Leu 86

					agc Ser 92											341
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					gat Asp 124											437
					aaa Lys 140											485
					aag Lys 156											533
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					gca Ala 204		tag *	ctgg	cagt	ca g	gaga	.ccgc	c go	tggg	gacg	683
tagt	tcca	tg g	aaag	tgac	c tc	tctg	gcac	ctg	gttt	tct	cttg	ttac	ac a	attg	aagaa	743
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Val Gly Ser Asp Pro Val Ile Leu Ala Thr Ala Gly Tyr Asp His Thr

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cac His 40	Gln	gac Asp	tcc Ser	cag Gln	gtg Val 45	Asn	gcc Ala	ttg Leu	gag Glu	gto Val 50	Thr	ccg Pro	gad Asp	c cgc	agc Ser 55	256
	Ile					Gln					Gly				atc Ile 71	304
cgc Arg 72	atg Met	tat Tyr	gat Asp	ctc Leu	aac Asn 77	tcc Ser	aat Asn	aac Asn	cct Pro	aac Asn 82	Pro	atc Ile	ato	ago Ser	tac Tyr 87	352
gac Asp 88	ggc	gtc Val	aac Asn	aag Lys	aac Asn 93	atc Ile	gcg Ala	tct Ser	gtg Val	ggc Gly 98	Phe	cac His	gaa Glu	gac	ggc Gly 103	400
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					tgc Cys 141											544
					gct Ala 157											592
					cct Pro 173											640
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ctc Leu 216	atc Ile	ccc Pro	aag Lys	act Thr	aag Lys 221	atc Ile	cct Pro	gcc Ala	cac His	acg Thr 226	cgc Arg	tac Tyr	gcc Ala	ctg Leu	cag Gln 231	784
					gac Asp 237											832

Gln Thr Cys Lys Ile Trp Arg Thr Ser Asn Phe Ser Leu Met Thr Glu 248 253 258 263	880
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•	CCa	cac	c af Mo	tg a et A 1	ac a sn T	cc to hr So	cc co er P:	ca go ro Gi 5	gc a ly T	cg g hr V	tg g al G	ly s	gt g er A 10	ac c sp P	cg g ro V	tc atc al Ile	169
	ctg Leu 15	Ala	c act	t gca r Ala	a ggo a Gly	tac Type	c Ası	c cac	c acc	gte r Vai	g cg l Ar	g Ph	c tg e Tr	g ca p Gl:	g gc n Al	c cac a His 30	217
	ago Ser 31	Gl)	ato / Ile	c tgo	c acc	c cgg Arg	y Thi	g gtg Val	g caq l Glr	g cad n His	c cag s Gl: 4:	n Asj	c tc p. Se	c cag r Gli	g gte n Val	g aat l Asn 46	265
	gcc Ala 47	Let	g gag ı Glu	g gto 1 Val	aca Thr	ccc Pro	Asp	cgc Arg	ago g Ser	ato Met	ati Ile 5	e Ala	t gc a Ala	t gca a Ala	a ggt a Gly	t tac y Tyr 62	313
ւնոք Ռույ Կւսո լեստ Գուս Առմ կոմ	cag Gln 63	cac	ato	cgc Arg	ato Met	tat Tyr 68	Asp	cto Leu	aac Asr	tco Ser	aat Asr 73	n Ası	c cci	aac Asr	c cco	atc Ile 78	361
ting from the	atc Ile 79	agc Ser	tac Tyr	gac Asp	ggc Gly	gto Val 84	Asn	aag Lys	aac Asn	ato Ile	gcg Ala 89	Ser	gtg Val	g ggo	tto Phe	cac His 94	409
dust 15. 454	gaa Glu 95	gac Asp	ggc Gly	cgc Arg	tgg Trp	atg Met 100	Tyr	acg Thr	ggc	ggc	gag Glu 105	Asp	tgo Cys	aca Thr	gcc Ala	agg Arg 110	457
	atc Ile 111	tgg Trp	gac Asp	ctc Leu	agg Arg	tcc Ser 116	cgg Arg	aac Asn	ctg Leu	cag Gln	tgc Cys 121	Gln	cgg Arg	atc Ile	ttc Phe	cag Gln 126	505
	gtg Val 127	aac Asn	gca Ala	ccc Pro	att Ile	aac Asn 132	tgc Cys	gtg Val	tgc Cys	ctg Leu	cac His 137	Pro	aac Asn	cag Gln	gca Ala	gag Glu 142	553
	ctc Leu 143	atc Ile	gtg Val	ggt Gly	gac Asp	cag Gln 148	agc Ser	gly ggg	gct Ala	atc Ile	cac His 153	Ile	tgg Trp	gac Asp	ttg Leu	aaa Lys 158	601
	aca Thr 159	gac Asp	cac His	aac Asn	gag Glu	cag Gln 164	ctg Leu	atc Ile	cct Pro	gag Glu	ccc Pro 169	gag Glu	gtc Val	tcc Ser	atc Ile	acg Thr 174	649
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	acc Thr 191	ctg Leu	gtg Val	ccc Pro	ttc Phe	tcc Ser 196	tgt Cys	cta Leu	ctt Leu	ccg Pro	ttg Leu 201	gcc Ala	atc Ile	ggc Gly	atc Ile	ctc Leu 206	745
	cag Gln	gaa Glu	gga Gly	gaa Glu	ttt Phe	gag Glu	tcc Ser	ttg Leu	gcc Ala	agg Arg	cgt Arg	ggc Gly	ctc Leu	ctg Leu	ttc Phe	ttg Leu	793

207	212	217	222
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gac gag gtg acc cag Asp Glu Val Thr Gln 239	ctc atc ccc aag Leu Ile Pro Lys 244	act aag atc cct gcc cac Thr Lys Ile Pro Ala His 249	c acg 889 s Thr 254
Arg Tyr Ala Leu Gln	tgt cgc ttc agc Cys Arg Phe Ser 260	ccc gac tcc acg ctc ctc Pro Asp Ser Thr Leu Leu 265	e gcc 937 1 Ala 270
Thr Cys Ser Ala Asp	cag acg tgc aag Gln Thr Cys Lys 276	atc tgg agg acg tcc aac Ile Trp Arg Thr Ser Asr 281	ttc 985 Phe 286
Ser Leu Met Thr Glu	ctg agc atc aag a Leu Ser Ile Lys 8 292	agc ggc aac ccc ggg gag Ser Gly Asn Pro Gly Glu 297	tcc 1033 Ser 302
Ser Arg Gly Trp Met	tgg ggc tgc gcc 1 Trp Gly Cys Ala 1 308	ttc tcg ggg gac tcc cag Phe Ser Gly Asp Ser Gln 313	tac 1081 Tyr 318
Ile Val Thr Ala Ser	tcg gac aac ctg g Ser Asp Asn Leu <i>1</i> 324	gcc cgg ctc tgg tgt gtg Ala Arg Leu Trp Cys Val 329	gag 1129 Glu 334
Thr Gly Glu Ile Lys	aga gag tat ggc g Arg Glu Tyr Gly 0 340	ggc cac cag aag gct gtt Bly His Gln Lys Ala Val 345	gtc 1177 Val 350
Cys Leu Ala Phe Asn A	gac agt gtg ctg g Asp Ser Val Leu G 356	gc tag cetgtgae ceeteg lly * 361	ggac 1228
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Asp Pro Asn Pro Ser Ala Ala Pro Thr Ser Thr Cys Ala Pro Arg Lys
                     19
                                          24
atg ccc aaa agg att tca ata tcc aaa caa ctg gct tca gtg aaa gct
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Met Pro Lys Arg Ile Ser Ile Ser Lys Gln Leu Ala Ser Val Lys Ala
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ctg agg aag tgc tca gat ctg gaa aaa gct att gcc acc act gct ctg
                                                                      194
Leu Arg Lys Cys Ser Asp Leu Glu Lys Ala Ile Ala Thr Thr Ala Leu
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att ttc aga aat tct tct gac tct gat ggt aaa ctt gaa aaa gct att
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Ile Phe Arg Asn Ser Ser Asp Ser Asp Gly Lys Leu Glu Lys Ala Ile
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                                                                      290
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Ala Lys Asp Leu Leu Gln Thr Gln Phe Arg Asn Phe Ala Glu Gly Gln
gaa acc aag cca aaa tac aga gag atc ctt tct gaa ctt gat gag cac
                                                                      338
Glu Thr Lys Pro Lys Tyr Arg Glu Ile Leu Ser Glu Leu Asp Glu His
                     99
aca gaa aat aag cta gat ttt gaa gac ttc atg atc ttg ctc tta agc
                                                                      386
Thr Glu Asn Lys Leu Asp Phe Glu Asp Phe Met Ile Leu Leu Ser
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                                         120
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                                                                      434
Ile Thr Val Met Ser Asp Leu Leu Gln Asn Ile Arg Asn Val Lys Ile
126
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                                         136
                                                             141
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Met Lys *
142
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			tat Tyr													400	
		_	gtg Val	_	-								_		-	448	
			ctt Leu													496	
			aca Thr													544	
			aaa Lys													592	
			cac His													640	
			aga Arg													688	
			tcc Ser													736	

ctg Leu 135	Asp	gaa Glu	gaa Glu	gcg Ala	cgg Arg 140	Pro	tca Ser	cac His	ato	cca Pro 145	Ala	ggt Gly	gac Asp	tgo Cys	agt Ser 150	784
gag Glu 151	His	tgg Trp	aag Lys	acg Thr	caa Gln 156	Arg	agt Ser	gag Glu	gag Glu	tat Tyr 161	Glu	gcg Ala	gaa Glu	ggc Gly	cag Gln 166	832
	Arg														tct Ser 182	880
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	atc Ile															1024
	gaa Glu															1072
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at Il 37	e Ar	g ca g Hi	t tt s Le	g ttg u Le	g gaa u Gli 380	ı Arç	a gti g Va:	t gai	t cag	g gte n Vai 389	l Va	c ag	a ga g Gl	a aa u Ly	a gag s Glu 390	1504
аа; Ly : 39:	s Le	a cg u Ar	g tc g Se	a gat r Ası	t cto Dei 396	ı Asp	aag Lys	g gco s Ala	gag a Glu	g aag 1 Lys 401	s Le	c aag	g tc s Se:	t tt r Le	a atg u Met 406	1552
gco Ala 401	a Se	g gag r Gl	g gtg u Val	g gat l Asp	gat Asp 412	His	cat His	gcg Ala	g gco Ala	ata Ile 417	e Glu	g; cgg	g cgg	g aa g As	t gag n Glu 422	1600
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ьет 439	ı Lys	s Asr	ı Glu	ı Leu	Arg 444	Lys	Glu	. Arg	Glu	Gln 449	Ile	Leu	Gln	Gli	g gca n Ala 454	1696
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G1u 471	Glu	Asn	Tyr	Ile	Arg 476	Asp	Arg	Leu	Ala	Leu 481	Ser	Leu	Lys	Glu	aac Asn 486	1792
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503	Glu	Asn	Leu	Thr	aac Asn 508	Lys	Leu	Gln	Arg	Asn 513	Leu	Glu	Asn	Val	Leu 518	1888
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cto Leu 631	ı Arç	a cte	g gag u Gli	g cto	gaa u Gli 630	ı Asp	aaa Lys	a gtg s Val	g cgo L Arg	cat His 641	ту:	t gaa r Gli	a aaq ı Lys	g cag s Gli	g ctg n Leu 646	2272
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				His	cat His 1388				Glu					Asn		4528
				Asn	aca Thr 1404				Glu					His		4576
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				Tyr	caa Gln .484				Leu					Lys		4816
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					Thr	cct Pro 1676				Gln					Leu		5392
					Lys	atg Met 1692				His					Leu		5440
					Lys	atc Ile .708				Leu					Lys		5488
						gct Ala											5536

gat gag cag ctc atg gag atg cag cac ctg agg tcc act gcg acg cct Asp Glu Gln Leu Met Glu Met Gln His Leu Arg Ser Thr Ala Thr Pro 1959 1964 1969 1974	6256
agc ccg tcc cct cat gct tgg gat ttg cag ctg ctc cag cag caa gcg Ser Pro Ser Pro His Ala Trp Asp Leu Gln Leu Leu Gln Gln Gln Ala 1975 1980 1985 1990	6304
tgc cgg atg gtg ccc agg gag cag ttt ttg cgg ttt aaa ccc cag ttg Cys Arg Met Val Pro Arg Glu Gln Phe Leu Arg Phe Lys Pro Gln Leu 1991 1996 2001 2006	6352
ctg cag gca gaa agg ata agc cag cac ttg cag ggg gaa ttt gaa aac Leu Gln Ala Glu Arg Ile Ser Gln His Leu Gln Gly Glu Phe Glu Asn 2007 2012 2017 2022	6400
agg acc tcc gaa acc aac aca cca cgg gga aac cgg gaa caa ttg gta Arg Thr Ser Glu Thr Asn Thr Pro Arg Gly Asn Arg Glu Gln Leu Val 2023 2028 2033 2038	6448
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tcc gtg ctg agc acc gac agc ggc aaa tcg gcg ccc gcc tct gcc acc Ser Val Leu Ser Thr Asp Ser Gly Lys Ser Ala Pro Ala Ser Ala Thr 3 8 13 18	164
gcg cgg gcc ctg gag cgc agg agg gac ccg gag ttg ccc gtc acg tcc Ala Arg Ala Leu Glu Arg Arg Arg Asp Pro Glu Leu Pro Val Thr Ser 19 24 29 34	212

	Asp					Leu					Glr				acc Thr 50	260
	Cys					Cys					Ala	acc Thr				308
	Asn											ctt Leu				356
												tca Ser				404
												gaa Glu				452
												gga Gly				500
												ttt Phe				548
gaa Glu 147	ctg Leu	tat Tyr	gaa Glu	gac Asp	agc Ser 152	ttg Leu	ctg Leu	gat Asp	cat His	tgt Cys 157	att Ile	act Thr	cat His	cac His	aga Arg 162	596
tcg Ser 163	gaa Glu	cgg Arg	agg Arg	cct Pro	gtg Val 168	ttc Phe	tgt Cys	cca Pro	ctt Leu	tgc Cys 173	cgt Arg	tta Leu	ata Ile	ccc Pro	gat Asp 178	644
												cat His				692
												aat Asn				740
									Arg			ctt Leu		Tyr		788
	cac His					taa *	tt t	tatt	aaaa	c ga	aggg	aaaa	999	acca	ctg	841
aatt	gcac	ca t	ttaa	gatg	c tg	cttg	aaca	aat	ggga	ggg	aagt	tgtc	aa t	gatt	gatgg	901
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Val 84	Glu	Phe	Gly	Asn	Gln 89	Leu	Glu	Gly	Lys	Trp 94	Ala	Val	Leu	Gly	Thr 99	
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									atc Ile							558
									atg Met							606
gtg Val 148	tag *	ttct	ctga	igc a	ıagtç	ıtggg	gg ca	agco	cagt	cgt	tcag	gaat	tgga	ıgcc <u>c</u>	jag	662
ttca	cgga	ıga o	agag	aacc	a ga	caga	caga	aga	ccca	gag	ccct	gggc	ca c	tcca	ctcct	722
gato	attt	ag c	tgcc	ggto	c ca	ctct	gacc	ttt	tgga	aag	aggo	tgtg	tg a	ggaa	.ggagt	782
agco	tggt	tg g	ıgggt	ctca	c tg	gcct	gact	ctg	caag	gaa	gaigg	tggc	tg c	actt	cccc	842
agct	tcca	gc t	ccag	acct	t ta	agco	ccag	gtg	cttg	, tgc	ctan	gatt	ta a	tgat	caaaa	902
aaaa	aaaa					•										910
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		1> C	DS 384)	(1	832)											
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agtt	gtcc	ct g	tcac	ttgg	t ga	gtgg	tctc	ttg	ttgc	ctg a	atati	tttaa	aa a	cgtt	tgata	120
tcat	ttca	aa g	ctat	agaa	a aa	ttgc	aaaa	aca	gtata	aag a	aaati	tata	at a	tacc	cttta	180
ccca	aatt	tg c	caag	tgtti	t ac	attt	ttcc	tgt	ttgct	tt (gttga	atgad	cc a	tcaa	ccaaa	240
tacc	acca	gg a	acat	acct	g ta	acca	gaga	cag	ctgat	tg (gtago	cttt	t g	cagc	aagga	300
agct	caca	ta c	cacg	gcaca	a at	tctg	cttt	ccci	tggaa	act 9	gtgto	catto	ca to	gact	ctgca	360
aatt	ccct	aa a	gtag	gagga	a aa	a.			atg Met							410

				gtg Val 15							458
				aag Lys 31							506
				gtg Val 47							554
				gaa Glu 63							602
				gga Gly 79							650
				cat His 95							698
				tta Leu 111							746
				cag Gln 127							794
		_		aca Thr 143	 _				 		842
				gat Asp 159							890
				tct Ser 175							938
-			_	ctt Leu 191		_	_	_	_		986
				gac Asp 207							1034
				cag Gln 223							1082

-			-					_	_	_	Ser	_			gtt Val 249	1130
	_				_	gga Gly						_		_	_	1178
						gat Asp										1226
						ttc Phe										1274
				_		aat Asn			_	_	_					1322
	_				_	aag Lys			_				_			1370
_	_		_		_	atg Met			_	-:-	_					1418
						agc Ser										1466
		-	_			gga Gly	_	_				_		-		1514
	_					aag Lys		_		_	_					1562
				_		tat Tyr		-	_	-	_		_	_		1610
						ttg Leu										1658
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						aaa Lys			_		_					1754
tgt	gaa	gac	tgt	ggg	aag	cgc	tac	aag	agg	cgc	ttg	aat	ctt	gat	ata	1802

Cys Glu Asp Cys Gly Lys Arg Tyr Lys Arg Arg Leu Asn Leu Asp Ile 458 463 468 473	
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cggaaagcaa aggc atg att tca cct tca ctt gaa ctg ctt cat tca gga Met Ile Ser Pro Ser Leu Glu Leu Leu His Ser Gly 1 5 10	170
ctc tgc aaa ttc cct gaa gta gaa gga aaa atg acc aca ttc aaa gag Leu Cys Lys Phe Pro Glu Val Glu Gly Lys Met Thr Thr Phe Lys Glu 13 18 23 28	218
gca gtg aca ttc aag gat gtg gct gtg gtc ttc act gag gag gag ctg Ala Val Thr Phe Lys Asp Val Ala Val Val Phe Thr Glu Glu Glu Leu 29 34 39 44	266
ggg ctg ctg gac cct gcc cag agg aag ctg tac cga gat gtg atg cta Gly Leu Leu Asp Pro Ala Gln Arg Lys Leu Tyr Arg Asp Val Met Leu 45 50 55 60	314
gag aac ttc agg aac ctg ctc tca gta ggg aat caa cca ttc cac caa Glu Asn Phe Arg Asn Leu Leu Ser Val Gly Asn Gln Pro Phe His Gln 61 66 71 76	362
gat act ttc cac ttc tta ggg aag gaa aag ttt tgg aag atg aag aca	410
Asp Thr Phe His Phe Leu Gly Lys Glu Lys Phe Trp Lys Met Lys Thr 77 82 87 92	
Asp Thr Phe His Phe Leu Gly Lys Glu Lys Phe Trp Lys Met Lys Thr	458

	Trp					Ser					, Ser				ata Ile 140	554
agg Arg 141	Asn	agc Ser	tct Ser	cag Gln	ttc Phe 146	Phe	aaa Lys	gaa Glu	ggt Gly	gat Asp 151	Val	cco Pro	tgc Cys	cag Glr	att Ile 156	602
											Lys				tgt Cys 172	650
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Lys 205	agc Ser	Phe	Cys	Tyr	Ser 210	Pro	Ala	Leu	His	Ile 215	His	Gln	Arg	Val	His 220	794
atg Met 221	gga Gly	gaa Glu	aaa Lys	tgc Cys	tat Tyr 226	aag Lys	tgt Cys	gat Asp	gtg Val	tgt Cys 231	ggt Gly	aag Lys	gaa Glu	ttt Phe	aat Asn 236	842
Gln 237	agc Ser	Ser	His	Leu	Gln 242	Thr	His	Gln	Arg	Val 247	His	Thr	Gly	Glu	Lys 252	890
Pro 253	ttc Phe	Lys	Arg	Gly	Gln 258	Cys	Gly	Lys	Gly	Phe 263	His	Ser	Arg	Ser	Ala 268	938
Leu 269	aat Asn	Val	His	Cys	Lys 274	Leu	His	Thr	Gly	Glu 279	Lys	Pro	Tyr	Asn	Cys 284	986
Glu 285	gaa Glu	Cys	Gly	Lys	Ala 290	Phe	Ile	His	Asp	Ser 295	Gln	Leu	Gln	Glu	His 300	1034
Gln 301	aga Arg	Ile	His	Thr	Gly 306	Glu	Lys	Pro	Phe	Lys 311	Cys	Asp	Ile	Cys	Gly 316	1082
aag Lys 317	agc Ser	ttc Phe	cgt Arg	gtt Val	aga Arg 322	tca Ser	aga Arg	ctt Leu	aat Asn	agg Arg 327	cat His	tcc Ser	atg Met	gtt Val	cac His 332	1130
aca Thr 333	gga Gly	gaa Glu	aaa Lys	cca Pro	ttc Phe 338	aga Arg	tgt Cys /	ġat Asp	Thr	tgt Cys 343	ggc Gly	aag Lys	aac Asn	ttt Phe	cgt Arg 348	1178
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Gln 349		g Ser	Ala	Let	Asr 354		His	s Sei	. Met	Va: 359		s Ile	e Glı	ı Glı	1 Lys 364	
	Туг					Суз					e Ile				gat Asp 380	1274
	Cys					Val					ı Lys				tgt Cys 396	1322
	Glu					Phe					Cyş	Leu			cat His 412	1370
											Cys				999 Gly 428	1418
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agg Arg 461	agg Arg	ttg Leu	gat Asp	ctt Leu	gag Glu 466	ttt Phe	cac His	cag Gln	agg Arg	gtc Val 471	cac His	acg Thr	ggt Gly	gag Glu	aga Arg 476	1562
ccc Pro 477	tat Tyr	aat Asn	tgt Cys	aag Lys	gaa Glu 482	tgt Cys	ggc Gly	aag Lys	agc Ser	ttt Phe 487	ggc Gly	tgg Trp	gcc Ala	tcc Ser	tgt Cys 492	1610
ctt Leu 493	ttg Leu	aaa Lys	cat His	cag Gln	aga Arg 498	ctc Leu	cac His	agt Ser	gly ggg	gaa Glu 503	aaa Lys	cct Pro	ttc Phe	aaa Lys	tgt Cys 508	1658
gaa Glu 509	gag Glu	tgt Cys	gga Gly	aag Lys	aga Arg 514	ttt Phe	act Thr	cag Gln	agt Ser	aca Thr 519	caa Gln	ctt Leu	cat His	tcc Ser	cat His 524	1706
cag Gln 525	aca Thr	tgc Cys	cat His	act Thr	gga Gly 530	gaa Glu	aag Lys	cta Leu	tac Tyr	aaa Lys 535	tgt Cys	gag Glu	cag Gln	tgt Cys	gag Glu 540	1754
aag Lys 541	gly aaa	tac Tyr	aac Asn	agt Ser	aaa Lys 546	ttt Phe	aat Asn	ctt Leu	gac Asp	atg Met 551	cac His	cag Gln	agg Arg	gtc Val	cac His 556	1802
agg Arg 557	gga Gly	gag Glu	cga Arg	ccc Pro	tat Tyr 562	aat Asn	tgt Cys	aag Lys	gaa Glu	tgt Cys 567	gga Gly	aag Lys	agc Ser	ttt Phe	ggc Gly 572	1850
tgg Trp	gct Ala	tca Ser	tgt Cys	ctt Leu	ttg Leu	aaa Lys	cat His	cag Gln	aga Arg	ctc Leu	cac His	agt Ser	gga Gly	gaa Glu	aag Lys	1898

573	578	583	588
cca ttg aaa tct g Pro Leu Lys Ser (589	gga gtg tgg gaa gag atc Gly Val Trp Glu Glu Ile 594	tac tca gaa ttc aca Tyr Ser Glu Phe Thr 599	gct 1946 Ala 604
tca ttt aca tca o Ser Phe Thr Ser V 605	gta agt cta tgt ggg aga Val Ser Leu Cys Gly Arg 610	aaa gcc ata taa Lys Ala Ile * 615	1988
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cccagcagtt cccttcg	gtgc gcggggggcg gcgagggt	ct tcagcagtcg ggagag	gccc 240
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ggc cca ggg ggc go Gly Pro Gly Gly Al 14	cc tcc tgc tcc gtg ggg g la Ser Cys Ser Val Gly A 19	la Pro Gly Gly Val S	cc 338 er 29
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gtg gat gtg gat ct Val Asp Val Asp Le 46	eg ctc ctg gga gag atc g eu Leu Leu Gly Glu Ile A 51	sp Pro Asp Gln Ala A	ac 434 sp 51
atc act tat gag gg Ile Thr Tyr Glu Gl 62	gg cga cag aag atg acc ag y Arg Gln Lys Met Thr Se 67	er Leu Ser Ser Cys Pl	tt 482 ne 77

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															caa Gln 109	578
					ttg Leu 115										cag Gln 125	626
					ctg Leu 131											674
					aag Lys 147											722
					aaa Lys 163											770
					gaa Glu 179											818
					gcg Ala 195											866
					caa Gln 211											914
		_		_	aag Lys 227						_	_	_			962
		_			act Thr 243					_	_					1010
					caa Gln 259											1058
					gtt Val 275										taa * 285	1106
ggaa	gato	at g	aagg	cctt	g go	attt	caat	tac	aggt	999	aaag	aaca	tg g	tgtt	ccaat	1166
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	Ser					Ser					Glu				aca Thr 112	695
	Val							_		-			_	_	gcg Ala 128	743
	Ala														CCC Pro 144	791
							ctg Leu								cac His 160	839
							gtg Val									887
_		_		_		_		_		_					ctg Leu 192	935
							cca Pro									983
						Asp	gag Glu									1031
gcc Ala 225	agg Arg	ata Ile	ctg Leu	gag Glu	tcc Ser 230	atc Ile	aca Thr	gtg Val	agc Ser	tcc Ser 235	ctg Leu	atg Met	gct Ala	aca Thr	ccc Pro 240	1079
							gtc Val									1127
							cct Pro									1175
							cgg Arg									1223
							tct Ser									1271

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cag gct cag atg gag gct ttc tga gggatcatct agcaaggagc tgtccagaga Gln Ala Gln Met Glu Ala Phe * 369 374	1517
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	gac tac tcc tac tgg gcg cac acc tca cct gag gac atc aac tac gcg Asp Tyr Ser Tyr Trp Ala His Thr Ser Pro Glu Asp Ile Asn Tyr Ala 53 68	365
	tcg caa aag cag gtg tac cgg gac atc ggc gag gag atg ctg cag cat Ser Gln Lys Gln Val Tyr Arg Asp Ile Gly Glu Glu Met Leu Gln His 69 74 79 \$	413
	gcc ttt gag gga tac aac gtg tgc atc ttc gcc tat ggg cag acg ggt Ala Phe Glu Gly Tyr Asn Val Cys Ile Phe Ala Tyr Gly Gln Thr Gly 85 90 95 100	461
4 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	gcc ggc aag tcc tac acc atg atg ggc aag cag gag aag gac cag cag Ala Gly Lys Ser Tyr Thr Met Met Gly Lys Gln Glu Lys Asp Gln Gln 101 106 111 116	509
free from from Som	ggc atc atc cca cag ctc tgc gag gac ctc ttc tct cgg atc aac gac Gly Ile Ile Pro Gln Leu Cys Glu Asp Leu Phe Ser Arg Ile Asn Asp 117 122 127 132	557
17	acg acc aac gac aac atg tcc tac tcc gtg gag gtc agc tac atg gag Thr Thr Asn Asp Asn Met Ser Tyr Ser Val Glu Val Ser Tyr Met Glu 133 148	605
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	att tac tgt gag cgc gtc cgt gac ctc ctg aac ccc aag aac aag ggc Ile Tyr Cys Glu Arg Val Arg Asp Leu Leu Asn Pro Lys Asn Lys Gly 149 154 159 164	653
17.4	aac ctt cgc gtg agg gag cac cca ctg ctg ggg ccc tac gtg gag gac Asn Leu Arg Val Arg Glu His Pro Leu Leu Gly Pro Tyr Val Glu Asp 165 170 175 180	701
	ctc tcc aag ctg gct gtc acc tcc tac aat gac atc cag gac ctc atg Leu Ser Lys Leu Ala Val Thr Ser Tyr Asn Asp Ile Gln Asp Leu Met 181 186 191 196	749
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	acc agc agt cgc tcc cac gcc gtc ttc aac atc atc ttc acc cag aag Thr Ser Ser Arg Ser His Ala Val Phe Asn Ile Ile Phe Thr Gln Lys 213 228	845
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acc ctg ggc aag g Thr Leu Gly Lys v 277	gtc atc tcc gcc Val Ile Ser Ala 282	c ctg gct gaa at a Leu Ala Glu Me 287	g gac tcc gga t Asp Ser Gly	ccc 1037 Pro 292
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Asn Lys Asn Lys I	ys Lys Lys Lys	Thr Asp Phe Ile		Asp
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Ser Val Leu Thr T	rp Leu Leu Arg	Glu Asn Leu Gly		Arg
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aca gct atg gtg g	ca gcc ctg agt	cct gca gac ato	Asn Tyr Asp	gag 1181
Thr Ala Met Val A	la Ala Leu Ser	Pro Ala Asp Ile		Glu
325	330	335		340
acc ctt agc acg c Thr Leu Ser Thr L 341	eu Arg Tyr Ala 346	Asp Arg Ala Lys	Gln Ile Arg (Cys 356
aat gct gtc atc a Asn Ala Val Ile A 357	sn Glu Asp Pro 362	Asn Asn Lys Leu 367	Ile Arg Glu I	seu 372
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Lys Asp Glu Val Ti	nr Arg Leu Arg	Asp Leu Leu Tyr		eu
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Ser Ser Leu Ser Al	a Leu Ser Ser	Arg Ala Ala Ser		eu
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His Glu Arg Ile Le	u Phe Ala Pro (Gly Ser Glu Glu		rg
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ctg aag gaa aca ga	g aag atc ata q	gct gag ctc aat	gag acc tgg ga	lu
Leu Lys Glu Thr Gl	u Lys Ile Ile <i>I</i>	Ala Glu Leu Asn	Glu Thr Trp Gi	
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Glu Lys Leu Arg Ar	g Thr Glu Ala 1	Ile Arg Met Glu	Arg Glu Ala Le	
453	458	463	46	
ctg gcc gag atg gg	t gtg gcc atg a	agg gag gat ggc (ggc acc ttg gg	.у .
Leu Ala Glu Met Gl	y Val Ala Met A	Arg Glu Asp Gly (Gly Thr Leu Gl	
469	474	479	48	

gt. Va: 48:	I Ph	c tc e Se	t cco	c aaa o Lys	a aag s Lys 490	Thi	a cca	a cad	c cto	c gt u Va 49	l As	c ct n Le	g aa u Asi	c ga n Gl	g gac u Asp 500	1661
Pro 50:	o rei	g at u Me	g tct t Sei	gag Glu	g tgo ı Cys 506	Leu	cto Lei	c tac ı Tyr	tac Tyi	c ate r Ile 51:	е Гу	g ga s As	t ggg p Gly	g at Y Il	c acc e Thr 516	1709
aga Arg 517	g Va.	1 Gl	c ago y Aro	g gag g Glu	gat Asp 522	Gly	gag Glu	g ago 1 Aro	g cgg g Arg	g cag g Glr 527	ı Ası	c at	t gtt e Val	cte	g agt u Ser 532	1757
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GLY 581	Asn	. Arg	atc Ile	Ile	Met 586	Gly	Lys	Ser	His	Val 591	Phe	Arg	Phe	Thr	His 596	1949
Pro 597	GIu	Gln	gcc Ala	Arg	Gln 602	Glu	Arg	Glu	Arg	Thr 607	Pro	Cys	Ala	Glu	Thr 612	1997
Pro 613	Ala	Glu	cct Pro	Val	Asp 618	Trp	Ala	Phe	Ala	Gln 623	Arg	Glu	Leu	Leu	Glu 628	2,045
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ccc Pro 693	gag Glu	gat Asp	gaa Glu	Val	cag Gln 698	tgg Trp '	aca Thr	gag Glu	Arg	gag Glu 703	tgt Cys	gag Glu	ctg Leu	gcg Ala	ctc Leu 708	2285

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gtg gcc gtc cag ggc cat ctc agc cga tga a gaggcccctg attatggctc Val Ala Val Gln Gly His Leu Ser Arg * 885 890	2864									
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	ı Lys					Thr					Ala				a caa s Gln 169	593
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	Asn					Gly					Val				agt Ser 201	689
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Arg 378		His	His	Gly	Glu 383		Lys	Pro	Tyr	388 388	_	a Asp	Val	. Сув	Asn 393	
	Gln					Ser					His				cat His 409	1313
	Gly										Gly				gct Ala 425	1361
	Ala										Ala				gaa Glu 441	1409
				_	_			ggg ggg	_	-		_	_		agt Ser 457	1457
								cat His								1505
								att Ile								1553
								aga Arg								1601
								aat Asn								1649
								cta Leu								1697
	_	_	_	_	_			caa Gln		_				_		1745
								act Thr								1793
								ctg Leu								1841
								tca Ser	Thr							1889
								tta Leu		tga *	cttt	gtaa	gga	atat	.gga	1940

602 607 612

attgctaaga	tatcattggt	agcaaacatc	tctggtaagg	tgcatatatt	catattaaat	2000
tcccattcat	ttgagttgtg	accattattt	ttcattcact	gaagtaaaag	cacctgatgc	2060
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ctgaattcca	gaaagaattt	tctagatata	aacttaatgc	atactagaaa	tcacttggct	2840
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<211> 2817

<212> DNA

<213> Homo sapiens

<220>

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<222> (109)..(2817)

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ttgtgggtgt cccggagcca gagggccaat tcctgtcccc ccagcagc atg gca tcg

Met Ala Ser

1

tgt gct gaa ccc tct gag ccc tct gcc cca ctg ccc gcc ggg gtc cca Cys Ala Glu Pro Ser Glu Pro Ser Ala Pro Leu Pro Ala Gly Val Pro

~	t				2	,				14	4				19	
ecc Pro 20	Let	gag 1 Glu	g gac 1 Asp	tto Phe	gag Glu 25	ı Val	cto Lev	gat 1 Asp	ggg Gly	ggtt Val	l Gl	g ga u Asj	t gca o Ala	a cag a Gli	g ggt n Gly 35	213
gag Glu 36	ı Glü	g gaa 1 Glu	ı gag ı Glu	gag Glu	g gag 1 Glu 41	ı Glu	gag Glu	gag Glu	g gaa Glu	gaç Glu 46	ı Ası	gao Asp	c cto	g agt 1 Sei	t gag r Glu 51	263
ctg Leu 52	Pro	ccg Pro	ttg Leu	gag	g gag 1 Glu 57	Thr	Gly	gag Glu	ggc Gly	cgg Arg	y Arg	d GJ7	gcg Ala	g gag a Glu	g gag 1 Glu 67	309
gag Glu 68	Asp	ctg Leu	ggg Gly	gaa Glu	gag Glu 73	Glu	gag Glu	gag Glu	ggc	cag Gln 78	Ala	g cac His	ctg Lev	g gag ı Glu	gac Asp 83	357
tgg Trp 84	Arg	Gly	cag Gln	ctc Leu	agt Ser 89	Glu	ctt Leu	ctg Leu	aac Asn	gac Asp 94	Phe	gag Glu	gag Glu	g agg L Arg	ctt Leu 99	405
gaa Glu 100	gac Asp	cag Gln	gaa Glu	cgc Arg	ctg Leu 105	aaa Lys	ccc Pro	gaa Glu	gga Gly	caa Gln 110	Arg	gaa Glu	cac	tcc Ser	gag Glu 115	453
gag Glu 116	cca Pro	ggt Gly	att Ile	tgc Cys	gag Glu 121	gag Glu	cag Gln	aac Asn	cca Pro	gag Glu 126	gcc Ala	gag Glu	cgg Arg	cga Arg	aga Arg 131	501
gag Glu 132	cag Gln	cag Gln	gga Gly	agg Arg	agc Ser 137	ggg Gly	gat Asp	ttc Phe	cag Gln	999 Gly 142	gcc Ala	gat Asp	cgt Arg	cct Pro	999 Gly 147	549
ccc Pro 148	gag Glu	gaa Glu	aag Lys	aga Arg	gaa Glu 153	gaa Glu	Gly 999	gac Asp	acg Thr	gag Glu 158	cct Pro	ctc Leu	ctg Leu	aaa Lys	caa Gln 163	597
gag Glu 164	gjà aaa	ccg Pro	gtg Val	gaa Glu	gcc Ala 169	gcg Ala	cag Gln	cct Pro	ccg Pro	gtg Val 174	gag Glu	agg Arg	aaa Lys	gaa Glu	gcc Ala 179	645
gcc Ala 180	gcc Ala	ctt Leu	gaa Glu	caa Gln	ggc Gly 185	cgc Arg	aag Lys	gtg Val	gag Glu	gag Glu 190	ctg Leu	cgg Arg	tgg Trp	cag Gln	gag Glu 195	693
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tcc Ser 212	gga Gly	Gly 999	aag Lys	cag Gln	att Ile 217	ctc Leu	ttt Phe	ccc Pro	Lys	gtc Val 222	aac Asn	ctg Leu	agc Ser	ccc Pro	gtg Val 227	789
acg Thr 228	ccc Pro	gca Ala	aag Lys	gac Asp	acg Thr 233	Gly ggg	ctc Leu	acc Thr	Ala	gct Ala 238	ccc Pro	cag Gln	gaa Glu	cca Pro	aag Lys 243	837

Pro				cag Gln						885
Pro				gtc Val						933
				aag Lys						981
				gcg Ala						1029
				agg Arg						1077
				gcg Ala						1125
	-	_	_	aac Asn	_	 _	_	_		1173
				ggt Gly						1221
				cgg Arg						1269
	_	_		cag Gln			_		 -	1317
				gct Ala						1365
				gcc Ala						1413
				gag Glu						1461
				aaa Lys						1509

											gtg Val					1557
											aac Asn					1605
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											ttc Phe					1701
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						Lys					ctc Leu					1845
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											ccg Pro					1941
											aaa Lys					1989
											cag Gln					2037
											gta Val					2085
											gag Glu					2133
gac Asp 676	cag Gln	gag Glu	agc Ser	agt Ser	gac Asp 681	cgc Arg	cgg Arg	cca Pro	ccc Pro	tcg Ser/ 686	ccc Pro	cca Pro	ggc Gly	ccc Pro	gag Glu 691	2181
gaa	agg	aag	gga	cag	aag	agg	gac	gag	gag	gaa	gag	gcg	aca	gag	agg	2229

Glu 692		Lys	Gly	Gln	Lys 697	Arg	Asp	Glu	Glu	Glu 702		Ala	Thr	Glu	Arg 707	
	Pro										Gln				tct Ser 723	2277
	Thr										Met				aga Arg 739	2325
	Ser														cag Gln 755	2373
	Leu										aag Lys					2421
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											agc Ser					2517
											aag Lys					2565
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											tct Ser					2661
atc Ile 852	tcc Ser	gac Asp	tcg Ser	gct Ala	ccc Pro 857	cca Pro	gcg Ala `	ccg Pro	ctg Leu	gta Val 862	aaa Lys	gaa Glu	gtc Val	acc Thr	aag Lys 867	2709
agg Arg 868	ttt Phe	tcc Ser	acc Thr	ccg Pro	gat Asp 873	gct Ala	gcc Ala	ccc Pro	gtg Val	tca Ser 878	aca Thr	gaa Glu	cca Pro	gcc Ala	tgg Trp 883	2757
ctg Leu 884	gct Ala	ttg Leu	gcc Ala	aaa Lys	agg Arg 889	aaa Lys	gca Ala	aag Lys	gct Ala	tgg Trp 894	agc Ser	gac Asp	tgt Cys	cca Pro	cag Gln 899	2805
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           <222> (50)..(436)
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                                                                Met Leu
      tgt ttg cac aag agc cga att gca gat ttg cag gat gtc ctg aag gag
                                                                             103
      Cys Leu His Lys Ser Arg Ile Ala Asp Leu Gln Asp Val Leu Lys Glu
ļ-Ł
     ccc tca att gca ttg gaa aag ctg cgg gaa ctc agc ttt agt ggc atc
                                                                             151
     Pro Ser Ile Ala Leu Glu Lys Leu Arg Glu Leu Ser Phe Ser Gly Ile
      19
                           24
     ccc tgt gag ggc gga ctg cgg tgc ctc tgc tgg aag att ctc ttg aac
                                                                            199
     Pro Cys Glu Gly Gly Leu Arg Cys Leu Cys Trp Lys Ile Leu Leu Asn
      35
                           40
     tac ctt ccc ttg gag agg gcc tca tgg acc tcc atc ctg gcc aag cag
                                                                            247
     Tyr Leu Pro Leu Glu Arg Ala Ser Trp Thr Ser Ile Leu Ala Lys Gln
      51
                           56
                                                                    66
     agg gag ctg tat gcc cag ttc ctg agg gaa atg atc atc cag cct ggc
                                                                            295
     Arg Glu Leu Tyr Ala Gln Phe Leu Arg Glu Met Ile Ile Gln Pro Gly
      67
     att gcc aag gcc aac atg ggt gtg tcc agg gag gat gtg act ttt gag
                                                                            343
     Ile Ala Lys Ala Asn Met Gly Val Ser Arg Glu Asp Val Thr Phe Glu
    gac cat toa etc aac ecc aac ect gac age egg tgg aac aeg tac tte
                                                                            391
     Asp His Ser Leu Asn Pro Asn Pro Asp Ser Arg Trp Asn Thr Tyr Phe
                                              109
     aag gac aac gag gtg ctg ctg cag atc gac aaa gat gtc cgg tag g
                                                                            437
     Lys Asp Asn Glu Val Leu Leu Gln Ile Asp Lys Asp Val Arg
     115
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<213> Homo sapiens

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<221> CDS
<222> (1716)..(1856)

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cctactgtgg ttgatcttaa catggttatc aggagtttcc ttgaatttaa tctagccctt	1680
atctaaactt cagtttactt taatagaaaa gaaaa atg gtc tgg atg tca ctt Met Val Trp Met Ser Leu 1	1733
ggg ccc aaa atc ttt tgg tac att tta gtg ccc ttt ctt cac cca ttc Gly Pro Lys Ile Phe Trp Tyr Ile Leu Val Pro Phe Leu His Pro Phe 7 12 17 22	1781
aga agt act tct gca gct gga atg gtc ctt cag aga aca ctc agt tct Arg Ser Thr Ser Ala Ala Gly Met Val Leu Gln Arg Thr Leu Ser Ser 23 28 33	1829
tct aga aag gag aca ctt aaa ata taa ttgcc tctccctccc atccctaact Ser Arg Lys Glu Thr Leu Lys Ile * 39 44	1881
catacagcca gaaagtcagg ttgaaggatt gtggaggact ctgctcttga acttttctgc	1941
acggacagca ggaccaccag tgcatgtgtg actcagtgtg gcatttgtct gcagggcata	2001
tgtacaccaa cccatccctg cctcttttta ccagtgcatg gatcatttct ctctcttctg	2061
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tgtggatcaa tgcttttgtc atgctctacc acgcaactga tgccatccaa atactgtgat	2181
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agcaatgctc cacagcacat ttggtcagaa aatacaatat ggcctcaagg ggtgctggtg	2301
cggcacagca gatgtttata tagagccatg gggccttaca acgtggcagt gccttcagat	2361
gtatctcatg cccgctttta tttcttattt catcgaccat taaggctgtt aaatctgctc	2421
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<211> 826

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (529)..(795)

<400> 1088
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ctacctggac taaacatttg c	ttcgccccg tctgcctgaa	tcgaacgtgc acttggcagt	240
cttcccttgt ccatacccgt t	tcctgggga tcgctacgga	ccttaaaata ccagcaacag	300
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ctgcagcctc aacctcctgg g	ctcaagcag tctttctgcc	tcagagtgct gggattacag	420
acgggagcca ccacgccagg c	ctacgagat ttgacttgta	aagaattaat tatattgact	480
gagagagaag cccagaagag g	aagaagagg aaagaaaagg	agtcaggg atg gct ctt Met Ala Leu 1	537
aca cag gga cct ttg aca Thr Gln Gly Pro Leu Thr 4	ttc agg gat gta gcc Phe Arg Asp Val Ala 14	ata gaa ttc tct cag Ile Glu Phe Ser Gln 19	585
gag gag tgg aaa tcc ctg Glu Glu Trp Lys Ser Leu 20 25			633
gtg atg ttg gag aac tac Val Met Leu Glu Asn Tyr 36 41	agg aac ctg gtc ttc Arg Asn Leu Val Phe 46	ctg ggt aag gat aat Leu Gly Lys Asp Asn 51	681
ttt gct cta gaa gtt aag Phe Ala Leu Glu Val Lys 52 57	atc tgc cct cgt gta Ile Cys Pro Arg Val 62	ttt ttg tat ttt ctc Phe Leu Tyr Phe Leu 67	729
tgt tgt ctt tct tgg gag Cys Cys Leu Ser Trp Glu 68 73	ccc ttc cat tat ttg Pro Phe His Tyr Leu 78	act gag act gaa gcc Thr Glu Thr Glu Ala 83	777
ttg ttg act cac aaa taa Leu Leu Thr His Lys *	aaagcttcat aatgctgaa	aa aaaaaaaaaa a	826

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<211> 765
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89

84

<220> <221> CDS

<222> (65)..(595)

<212> DNA

<213> Homo sapiens

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ctg ctc ttc agc cac ctc tct gcg gtc cag acg agg ggc atc aag cac Leu Leu Phe Ser His Leu Ser Ala Val Gln Thr Arg Gly Ile Lys His 16 21 26 31	157
aga atc aag tgg aac cgg aag gcc ctg ccc agc act gcc cag atc act Arg Ile Lys Trp Asn Arg Lys Ala Leu Pro Ser Thr Ala Gln Ile Thr 32 37 42 47	205
gag gcc cag gtg gct gag aac cgc ccg gga gcc ttc atc aag caa ggc Glu Ala Gln Val Ala Glu Asn Arg Pro Gly Ala Phe Ile Lys Gln Gly 48 53 58 63	253 ·
cgc aag ctc gac att gac ttc gga gcc gag ggc aac agg tac tac gag Arg Lys Leu Asp Ile Asp Phe Gly Ala Glu Gly Asn Arg Tyr Tyr Glu 64 69 74 79	301
gcc aac tac tgg cag ttc ccc gat ggc atc cac tac aac ggc tgc tct Ala Asn Tyr Trp Gln Phe Pro Asp Gly Ile His Tyr Asn Gly Cys Ser 80 85 90 95	349
gag gct aat gtg acc aag gag gca ttt gtc acc ggc tgc atc aat gcc Glu Ala Asn Val Thr Lys Glu Ala Phe Val Thr Gly Cys Ile Asn Ala 96 101 106 111	397
acc cag gcg gcg aac cag ggg gag ttc cag aag cca gac aac aag ctc Thr Gln Ala Ala Asn Gln Gly Glu Phe Gln Lys Pro Asp Asn Lys Leu 112 117 122 127	445
cac cag cag gtg ctc tgg cgg ctg gtc cag gag ctc tgc tcc ctc aag His Gln Gln Val Leu Trp Arg Leu Val Gln Glu Leu Cys Ser Leu Lys 128 133 138 143	493
cat tgc gag ttt tgg ttg gag agg ggc gca gga ctt cgg gtc acc atg His Cys Glu Phe Trp Leu Glu Arg Gly Ala Gly Leu Arg Val Thr Met 144 149 154 159	541
cac cag cca gtg ctc ctc tgc ctt ctg gct ttg atc tgg ctc acg gtg His Gln Pro Val Leu Leu Cys Leu Leu Ala Leu Ile Trp Leu Thr Val 160 165 170 175	589
aaa taa gcttgccagg aggctggcag tacagagtgc agcagcgagc aaatcctggc Lys * 176	645
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cggcggcggc gctaccaagc cggcggtctc cggcaagcag ggcaatgtgc tcccgctct	g 180
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ctg tcg tcg cct tac ttc aaa gta cag ctc tac gag ctc aag acc tac Leu Ser Ser Pro Tyr Phe Lys Val Gln Leu Tyr Glu Leu Lys Thr Tyr 12 17 22 27	277
cac gag gtg gtg gac gag atc tac ttt aag gtc acg cac gtt gaa cca His Glu Val Val Asp Glu Ile Tyr Phe Lys Val Thr His Val Glu Pro 28 33 38 43	325
tgg gag aaa gga agc agg aaa aca gcg ggc cag aca ggg atg tgc gga Trp Glu Lys Gly Ser Arg Lys Thr Ala Gly Gln Thr Gly Met Cys Gly 44 49 54 59	373
ggg gtt cga ggt gtt gga aca gga gga att gtt tct aca gca ttt tgc Gly Val Arg Gly Val Gly Thr Gly Gly Ile Val Ser Thr Ala Phe Cys 60 65 70 75	421
ctg tta tac aaa tta ttt acc ctg aag tta act cga aag caa gtg atg Leu Leu Tyr Lys Leu Phe Thr Leu Lys Leu Thr Arg Lys Gln Val Met 76 81 86 91	469
ggt ctt ata aca cac aca gac tct cca tat att aga gcg ctt gga ttt Gly Leu Ile Thr His Thr Asp Ser Pro Tyr Ile Arg Ala Leu Gly Phe 92 97 102 107	517
atg tat ata aga tat aca cag ccc cct aca gat ctg tgg gac tgg ttt Met Tyr Ile Arg Tyr Thr Gln Pro Pro Thr Asp Leu Trp Asp Trp Phe 108 113 118 123	565
gaa tcc ttc ctt gat gat gaa gag gac cta gat gtg aag gct ggt gga Glu Ser Phe Leu Asp Asp Glu Glu Asp Leu Asp Val Lys Ala Gly Gly 124 129 134 139	613
ggc tgt gta atg acc att gga gaa atg cta cga tct ttt ctc aca aaa Gly Cys Val Met Thr Ile Gly Glu Met Leu Arg Ser Phe Leu Thr Lys 140 145 150 155	661

											cca Pro					709
_			_		_				_		aga Arg					75 <i>7.</i>
											cat His					805
_						_			_		cgg _į Arg					853
											cat His					901
	_	_	_		_	_				-	cag Gln					949
_	_		_		_		_		_	_	tcc Ser					997
			_	_			_		_		gaa Glu					1045
-	-	_	_	_	_				-	_	agg Arg			-		1093
_	_			-			_		_	_	cga Arg		_			1141
											aga Arg					1189
											gta Val					1237
		_	_		_	_				_	gat Asp	_				1285
											agc Ser					1333
cac	aga	agt	agg	agt	cga	agt	aga	aat	gca	999	aaa	cga	agt	aga	agt	1381

His Arg Ser Arg Ser Arg Ser Arg Asn Ala Gly Lys Arg Ser 380 385 390	Arg Ser
aga agc aaa gag aaa tca agt aaa cat aaa aat gaa agt aaa Arg Ser Lys Glu Lys Ser Ser Lys His Lys Asn Glu Ser Lys 396 401 406	
tca aat aaa cga agt cga agt ggc agt caa gga aga act gac Ser Asn Lys Arg Ser Arg Ser Gly Ser Gln Gly Arg Thr Asp 412 417 422	e agt gtt 1477 o Ser Val 427
gaa aaa tca aaa aaa cgg gaa cat agt ccc agc aaa gaa aaa Glu Lys Ser Lys Lys Arg Glu His Ser Pro Ser Lys Glu Lys 428 433 438	
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gaa caa gag agc caa gaa aaa cag cat aaa aac aaa gat gag Glu Gln Glu Ser Gln Glu Lys Gln His Lys Asn Lys Asp Glu 476 481 486	
tga aaat attttgtaaa agtggatcac attgaatcct ataaatgatt aa * 492	atctgctt 1726
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tttgtttttg aaatgtacag tctgtacata tgtcctgaaa atgttttaat	tcctttggca 1966
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ctggattctt cattactaca tctgaaaagc ttctc	catcta gaagteeteg accatgttta 180
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tcacatacat atagaactta gccctttctc ctgct	actgc tgctgcaaca actttgacct 600
	tt gtt gtg ata tat gat tta 653 The Val Val Ile Tyr Asp Leu 5
cga gat atg cca gcc aat cat cag aca tt Arg Asp Met Pro Ala Asn His Gln Thr Ph 9 14	
tct gta cct gtt aaa caa gaa gtg aag ag Ser Val Pro Val Lys Gln Glu Val Lys Ar 25 30	
atc cga cac aca gaa gaa cgg tta tta cg Ile Arg His Thr Glu Glu Arg Leu Leu Ar 41	
ttc cag agt tct aaa tct gga aag atc ta Phe Gln Ser Ser Lys Ser Gly Lys Ile Ty 57 62	
ctc ctg ttc tct aga aag tca atg gag gt Leu Leu Phe Ser Arg Lys Ser Met Glu Va 73 78	
gaa ctc aaa tct tac act gaa tca cca ac Glu Leu Lys Ser Tyr Thr Glu Ser Pro Th 89 94	
aga tgt tga taaggag tgatgattta aagtatt Arg Cys * 105	tac tcagtaccca agctgtgcga 997
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														cta Leu		687
gca Ala 124	gaa Glu	gta Val	gaa Glu	aaa Lys	tac Tyr 129	aaa Lys	gac Asp	tgt Cys	gat Asp	ccg Pro 134	caa Gln	gtt Val	gtg Val	gaa Glu	gaa Glu 139	735
ata Ile 140	cgc Arg	caa Gln	gca Ala	aat Asn	aaa Lys 145	gta Val	gcc Ala	aaa Lys	gaa Glu	gct Ala 150	gct; Ala	aac Asn	aga Arg	tgg Trp	act Thr 155	783
														ggg Gly		831
														ttt Phe		. 879
	ata Ile		taa *	aat	atto	cate	gt g	ggtga	aagga	at.gt	acaa	igctt	gtg	gaata	atgt	934
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atca	atttt	at t	taatg	gttaa	aa ta	aagt	gtaa	ı aat	gcag	gatg	ttct	tcac	cc c	ettt	ggtag	1054
aaca	aaag	gca g	ggatg	gataa	ac ca	tato	cccc	cas	gtgct	cat	caaa	gtag	ga c	cacta	aaaat	1114
ccat	ccat	ct d	cagto	caaag	gt c <u>c</u>	gagcg	gccg	g cga	attt	agt	agta	gtag	lcd d	gccgc	tctag	1174
agga	itcca	ag d	cttac	gtac	g cg	gtgca	tgcg	g acc	gtcat	agc	tctt	ctat	ag t	gtca	ıcctaa	1234
atto	caagt	t														1243

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		agaa	agg	aatt	ctag	ca a	agtt	atto	t gt	aaat	acaa	cag	aato	agt	tgct	tttgtt	. 1	180
	aaa	tata	ittg	gtgc	aagt	tc a	ac	Me								g ggt u Gly	2	230
	gtg Val 10	Tyr	cca Pro	tct Ser	tac Tyr	cca Pro 15	gac Asp	ctg Leu	gtc Val	atc Ile	gat Asp 20	Val	gga Gly	gaa Glu	gtg Val	act Thr 25	2	27 E
-		Gly					Lys					Thr				caa Gln 41	3	326
	gag Glu 42	Arg	gcg Ala	aga Arg	gtt Val	ata Ile 47	cgg Arg	gcc Ala	gcg Ala	tgt Cys	gct Ala 52	tta Leu	tta Leu	aac Asn	tca Ser	gga Gly 57	3	74
	gga Gly 58	Gly	gtg Val	att Ile	cag Gln	atg Met 63	gaa Glu	atg Met	gcc Ala	aac Asn	agg Arg 68	gat Asp	gag Glu	cgt Arg	ccc Pro	aca Thr 73	4	22
	gag Glu 74	atg Met	gga Gly	ctg Leu	gat Asp	tta Leu 79	gaa Glu	gaa Glu	tcc Ser	ttg Leu	aga Arg .84	aag Lys	ctt Leu	att	cag Gln	tat Tyr 89	4	70
	cca Pro 90	Tyr	ttg Leu	cag Gln	gct Ala	ttc Phe 95	ttt Phe	gag Glu	act Thr	aag Lys	caa Gln 100	cac His	gga Gly	agg Arg	tgt Cys	ttt Phe 105	· 5	18
	tat Tyr 106	att Ile	ttt Phe	gtt Val	aaa Lys	tct Ser 111	tgg Trp	agt Ser	ggt Gly	gat Asp	cct Pro 116	ttc Phe	ctt Leu	aaa Lys	gat Asp	ggt Gly 121	5	66
	tct Ser 122	ttc Phe	aat Asn	tcc Ser	cgc Arg	att Ile 127	tgc Cys	agc Ser	ctt Leu	act Thr	tcg Ser 132	gct Ala	atc Ile	tac Tyr	atg Met	cag Gln 137	6:	14
	atg Met 138	aat Asn	gag Glu	acc Thr	agg Arg	ccc Pro 143	tta Leu	ttt Phe	cac His	acc Thr	ata Ile 148	tac Tyr	taa *	aato	atct	ca	66	63
	aaat	acat	ta t	aatg	caga	c at	aaga	tgcg	aaa	.ctgt	aaa	acta	.ctgg	raa g	aaaa	cacag	. 72	23
	ggga	aaat	ct c	catg	atac	t gg	tctg	ggca	atg	tttt	cct	ggat	atga	.cc c	caaa	tacac	78	3
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ctgtttcaga gcctccacac c atg acc tcc aag aag ctg gtg aac tcg gtg Met Thr Ser Lys Leu Val Asn Ser Val 1 5																231		
	gct Ala 11	ggc Gly	tgt Cys	gct Ala	gat Asp	gac Asp 16	Ala	ctt Leu	gct Ala	ggc Gly	ctg Leu 21	gtg Val	gcc Ala	tgc Cys	aac Asn	ccc Pro 26		279
						cag Gln 32												327
	gac Asp 43	agc Ser	ctc Leu	aag Lys	ggc Gly	cgg Arg 48	gtg Val	gca Ala	·ctg Leu	ctģ Leu	tcg Ser ,53	ggt Gly	Gly 999	ggc Gly	tct Ser	ggc Gly 58		375
						gct Ala 64												423
						gtg Val 80											•	471
						gtg Val 96											. !	519
						act Thr 112											9	567
						gaa Glu 128											•	515
						act Thr 144				Lys							•	63
						ata Ile 160											-	711
						gag Glu 176											•	759

						gtg Val					Cys				ggt Gly 202	807
						ctc Leu										855
999 Gly 219	atc Ile	cac His	Gly 999	gaa Glu	gct Ala 224	ggt Gly	gtg Val	cgc Arg	cgg Arg	ata Ile 229	aag Lys	atg Met	gca Ala	acc Thr	gcc Ala 234	903
	Glu					atg Met										951
						cag Gln										999
						tca Ser										1047
						gag Glu										1095
						tca Ser										1143
						gag Glu										1191
						cct Pro										1239
						gcc Ala										1287
		_				tca Ser	_	_	_		_		_			1335
						ctc Leu										1383
/ ctg Leu 395	gac Asp	cgg Arg	gct Ala	gct Ala	ggt Gly 400	gac Asp	ggc Gly	gac Asp	tgt Cys	ggc Gly 405	acc Thr	acc Thr	cac His	agc Ser	cgt Arg 410	1431

gcg gcc aga gca atc cag gag tgg ctg aag gag ggc cca ccc cct gcc Ala Ala Arg Ala Ile Gln Glu Trp Leu Lys Glu Gly Pro Pro Pro Ala 411 416 421 426	1479
agc cct gcc cag ctg ctc tcc aag ttg tct gtt ctg ctc ctg gag aag Ser Pro Ala Gln Leu Leu Ser Lys Leu Ser Val Leu Leu Leu Glu Lys 427 432 437 442	1527
atg gga ggc tca tct ggg gcg ctc tat ggc ctg ttc ctg act gcg gct Met Gly Gly Ser Ser Gly Ala Leu Tyr Gly Leu Phe Leu Thr Ala Ala 443 458	1575
gca cag ccc ctg aag gcc aag acc agc ctc cca gcc tgg tct gcc Ala Gln Pro Leu Lys Ala Lys Thr Ser Leu Pro Ala Trp Ser Ala Ala 459 464 469 474	1623
atg gat gcc ggc ctg gaa gcc atg cag aag tat ggc aag gct gct cca Met Asp Ala Gly Leu Glu Ala Met Gln Lys Tyr Gly Lys Ala Ala Pro 475 480 485 490	1671
ggg gac agg act atg ctg gat tct ctg tgg gca gcg ggg cag gag ctc Gly Asp Arg Thr Met Leu Asp Ser Leu Trp Ala Ala Gly Gln Glu Leu 491 496 501 506	1719
caa gcc tgg aag agc cca gga gct gat ctg tta caa gtc ctg acc aaa Gln Ala Trp Lys Ser Pro Gly Ala Asp Leu Leu Gln Val Leu Thr Lys 507 512 517 522	1767
gca gtc aag agt gcc gaa gct gca gcc gag gcc acc aag aat atg gaa Ala Val Lys Ser Ala Glu Ala Ala Ala Glu Ala Thr Lys Asn Met Glu 523 528 533 538	1815
gct gga gcc gga aga gcc agt tat atc agc tca gca cgg ctg gag cag Ala Gly Ala Gly Arg Ala Ser Tyr Ile Ser Ser Ala Arg Leu Glu Gln 539 544 549 554	1863
cca gac ccc ggg gcg gtg gca gct gct gcc atc ctc cgg gcc atc ttg Pro Asp Pro Gly Ala Val Ala Ala Ala Ala Ile Leu Arg Ala Ile Leu 555 560 565 570	1911
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                                                                       165
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Met Trp Phe Met Tyr Leu Leu Ser Trp Leu Ser Leu Phe Ile Gln Val
gcc ttc atc acg ctg gct gtc gcg gct gga ctc tat tac ctg gca gaa
                                                                       213
Ala Phe Ile Thr Leu Ala Val Ala Ala Gly Leu Tyr Tyr Leu Ala Glu
                                          27
                                                                      261
ctg ata gaa gaa tac aca gtg gcc acc agc agg atc ata aaa tac atg
Leu Ile Glu Glu Tyr Thr Val Ala Thr Ser Arg Ile Ile Lys Tyr Met
                     38
                                          43
                                                                      309
atc tgg ttc tcc acc gct gta ctg att ggc ctc tac gtc ttt gag cgc
Ile Trp Phe Ser Thr Ala Val Leu Ile Gly Leu Tyr Val Phe Glu Arg
                                          59
                     54
                                                                      357
tte ecc acc age att gga gtg gge eta tte acc aac etc gte tac
Phe Pro Thr Ser Met Ile Gly Val Gly Leu Phe Thr Asn Leu Val Tyr
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                     70
 65
                                                                      405
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Phe Gly Leu Leu Gln Thr Phe Pro Phe Ile Met Leu Thr Ser Pro Asn
81
                     86
                                                                      453
ttc atc ctg tcg tgt gga cta gtg gtg gtg aat cat tac cta gca ttt
Phe Ile Leu Ser Cys Gly Leu Val Val Val Asn His Tyr Leu Ala Phe
97
                    102
                                                             112
                                                                      501
cag ttt ttt gca gaa gaa tat tat ccc ttc tca gag gtc ctg gcc tat
Gln Phe Phe Ala Glu Glu Tyr Tyr Pro Phe Ser Glu Val Leu Ala Tyr
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                                        123
ttc act ttc tgc ctg tgg ata att ccg ttt gcg ttt ttt gtg tca ctt
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Phe Thr Phe Cys Leu Trp Ile Ile Pro Phe Ala Phe Phe Val Ser Leu
                    134
                                        139
                                                                      597
tcg gcc ggg gag aac gtc ctg ccc tct acc atg cag cca gga gat gat
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145
                    150
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agg ctg gtc tcg gag acc gca gag cgg aaa cag gag cag aag a	157
acc gag gcg gag gct gcc gac tcc ggt gcc gtc gga gcc cgc cgc ttc. Thr Glu Ala Glu Ala Ala Asp Ser Gly Ala Val Gly Ala Arg Arg Phe 30 45	205
ctg ctc tgt ctc tac ttg ggg ggc ttc ttg gat ttg ttt ggt gtc agc Leu Leu Cys Leu Tyr Leu Gly Gly Phe Leu Asp Leu Phe Gly Val Ser	253

40					53	L				56	5				61	
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cca Pro 78	Thr	gtt Val	gct L Ala	gga Gly	ata 7 Ile 83	· Val	a ggo . Gly	tco Ser	tcc Ser	tat Tyr 88	Gly	c att y Ile	ttq E Le	g ca ı Gl	a ctc n Leu 93	349
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tct Ser 110	tcc Ser	ttg Leu	g ctg Leu	gca Ala	tgc Cys 115	Ile	cta Leu	ctc Leu	agt Ser	gct Ala 120	Lev	g ggc ı Gly	tat Tyr	cto Lev	c ctt 1 Leu 125	445
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gaa Glu 222	gca Ala	aaa Lys	ccg Pro	ggc Gly	agt Ser 227	aca Thr	gag Glu	aag Lys	ggc Gly	ctg Leu 232	cca Pro	ttg Leu	cga Arg	aag Lys	acc Thr 237	781
cat (His) 238	gtg Val	ctg Leu	ttg Leu	gga Gly	agg Arg 243	agc Ser	cat His	gac Asp	Thr	gtg Val 248	cag Gln	gag Glu	gca Ala	gcc Ala	acc Thr 253	829
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gta 9 Val 1 270	gtg Val	ttg Leu	gcc Ala	Leu	cgg Arg 275	aac Asn	atg Met	aag Lys	Asn	ctg Leu 280	ctg Leu	ttt Phe	tcc Ser	gaa Glu	atg Met 285	925

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	tac Tyr 302	tac Tyr	agt Ser	aac Asn	ttt Phe	gto Val 307	. Leu	gcc	ctg Leu	gag Glu	gag Glu 312	Arg	ttt Phe	gly ggg	gtg Val	cgg Arg 317	1021
	ccc Pro 318	aag Lys	gtg Val	aca Thr	ggc	tac Tyr 323	Leu	atc Ile	agt Ser	tac Tyr	ago Ser 328	Ser	atg Met	ctg Leu	ggg Gly	gcc Ala 333	1069
	gtg Val 334	gcc Ala	ggc	ctt Leu	gcc Ala	ctg Leu 339	gly aaa	cca Pro	atc Ile	cta Leu	cgg Arg 344	ctg Leu	tac Tyr	aag Lys	cac His	aac Asn 349	1117
•	tcg Ser 350	cag Ġln	gca Ala	ctg Leu	ctg Leu	ctg Leu 355	cat His	tcc Ser	agc Ser	ata Ile	ctc Leu 360	acc Thr	tgc Cys	aca Thr	ctg Leu	ctg Leu 365	1165
J	ctg Leu 366	ctc Leu	tac Tyr	tcc Ser	ttg Leu	gcc Ala 371	ccc Pro	acc Thr	atg Met	ggt Gly	gca Ala 376	gtt Val	gtc Val	ctc Leu	tcc Ser	tcc Ser 381	1213
•	act Thr 882	ctc Leu	ctg Leu	tcc Ser	ttc Phe	tcc Ser 387	act Thr	gcc Ala	att Ile	ggc Gly	agg Arg 392	acg Thr	tgc Cys	atc Ile	acg Thr	gac Asp 397	1261
1	etc eu 198	cag Gln	ctg Leu	act Thr	gtg Val	ggc Gly 403	Gly aaa	gcc Ala	cag Gln	gcc Ala	agc Ser 408	ggc Gly	acc Thr	ctt Leu	att Ile	ggc Gly 413	1309
V	tg al 14	gly 999	cag Gln	tct Ser	Val	act Thr 419	gca Ala	gtg Val	ggc Gly	Arg	atc Ile 424	atc Ile	gcc Ala	cct Pro	Leu	ctc Leu 429	1357
S	cg er 30	ggg Gly	gtt Val	gcc Ala	Gln	gag Glu 435	gtc Val	agc Ser	cct Pro	Cys	ggc Gly 440	ccc Pro	ccc (agc Ser	Leu	ggc Gly 445	1405
A	ct (la ' 46	gtg Val 1	tta Leu .	gcc i Ala i	Leu	gtg Val 451	gcc (Ala	att Ile	ttc Phe	Ile	atg Met 456	tct Ser	cta a Leu A	aac a Asn 1	Lys :	cga Arg 461	1453
н	ac 1 is 1 62	tct a Ser s	agt (Ser (ggt g Gly <i>I</i>	Asp (999 Gly 467	aat a Asn S	agt : Ser :	aaa 1 Lys 1	Leu 1	aaa a Lys : 472	agt (Ser (gag t Glu	ag a *	atgga	att	1502
t	ggad	caaca	at a	aagca	aca	a aa	tttga	agat	ggt	gaat	tga g	gggc	cggac	g co	atga	atgaa	1562
a	aggg	jcact	t to	ggaaa	agggi	t tg	gggtg	ggaa	9998	aaata	att 1	ccg	ggtgg	g to	gtgag	gctgt	1622
t	3999	ttcc	a g	gtcag	gctci	t tg	gccat	gca	gcca	atgco	ctg d	cagga	atgat	c ag	aagt	cacg	1682
g	caco	tcat	g gg	gaagg	gttaa	a ga	ctgga	igca	aago	tttt	cc a	agto	gtgac	с са	gaga	ccag	1742

tagcatcagc	atcacctgag	acctgtgaac	actgaagctc	cagctcagac	atgttgggga	1802
ccattttaat	aagataccta	gctgattttt	tgcacagcaa	actttgaaaa	cccctggtct	1862
aaggggtagt	atttgtatca	cttatggaat	ataatctcag	ggaaattaaa	ttctgctcaa	1922
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			aac Asn														554
			acc Thr														602
			act Thr														650
	ttc Phe 175	aaa Lys	ctg Leu	aag Lys	aag Lys	aga Arg 180	gaa Glu	aag Lys	agg Arg	cgc Arg	tct Ser 185	act Thr	agc Ser	cga Arg	cag Gln	ttt Phe 190	698
			ggt Gly														746
	gca Ala 207	gta Val	gac Asp	cag Gln	ggg Gly	aat Asn 212	gaa Glu	tcc Ser	ata Ile	gtt Val	gca Ala 217	aaa Lys	act Thr	aca Thr	gtg Val	act Thr 222	794
			aat Asn														842
			tat Tyr														890
			agt Ser														938
	_		gac Asp	_			_		_	_							986
		_	ttt Phe	_		_	-	_				_					1034
			aag Lys														1082
	-	_	gtg Val	_				_	_		_	_	_				1130
,	_		cct Pro		_					_	_		-				1178
	ctg	gca	gac	ttt	gtg	tcc	cag	act	tct	cca	atg	atc	ccc	tcc	att	gtt	- 1226

Leu 351		Asp	Phe	Val	Ser 356	Gln	Thr	Ser	Pro	Met 361		Pro	Ser	Ile	Val 366		
											Leu				ggc Gly 382	•	1274
_						_		_		_			_		gag Glu 398		1322
			_		aaa Lys 404		_				_			_	gat Asp 414		1370
		_		_	agc Ser 420				_			_					1418
_			_		ttt Phe 436	-			_	_	Phe	_	_	_			1466
_			_	-	gac Asp 452		_		_	_	_			_			1514
					gcc Ala 468												1562
					gct Ala 484												1610
					ttt Phe 500												1658
					aca Thr 516												1706
	Val				ctt Leu 532												1754
					gag Glu 548												1802
					aca Thr 564												1850
					act Thr												1898

575 586	0 58	5 590	
tct ggt tcc ctg tca cas Ser Gly Ser Leu Ser Gli 591 596	n Arg Val Arg Ser Th	r Leu Thr Lys Asn Thr	1946
cct aga ttt ggg agc aac Pro Arg Phe Gly Ser Lys 607 612	s Ser Lys Ser Ala Th	r Asn Leu Gly Arg Gln	1994
ggc aac ttt ttt gct tct Gly Asn Phe Phe Ala Ser 623	r Pro Met Leu Lys *	,	2045
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gtactcatta ctacttttag o	cattctccag gcttttact	c aagtttaatt gtgcatgagg	2165
gttttattaa aactatatat a	atctcccctt ccttctcct	c aagtcacata atatcagcac	2225
tttgtgctgg tcattgttgg g	gagcttttag atgagacat	c tttccagggg tagaagggtt	2285
agtatggaat tggttgtgat t	tettttggg gaaggggt	t attgttcctt tggcttaaag	2345
ccaaatgctg ctcatagaat g	gatctttctc tagtttcat	t tagaactgat ttccgtgaga	2405
caatgacaga aaccctacct a	atctgataag attagcttg	t ctcagggtgg gaagtgggag	2465
ggcagggcaa agaaaggatt a	agaccagagg atttaggat	g cctccttcta agaaccagaa	2525
gttctcattc cccattatga a	actgagctat aatatggag	c tttcataaaa atgggatgca	2585
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attcattctg atgcttggcc c	cccatacccc caaccttgt	c cagtggagcc caacttctaa	2825
aggtcaatat atcatccttt g	ggcatcccaa ctaacaataa	a agagtaggct ataagggaag	2885
attgtcaata ttttgtggta a	agaaaagcta cagtcattt	ttctttgcac tttggatgct	2945
gaaatttttc ccatggaaca t	agccacatc tagatagat	g tgagcttttt cttctgttaa	3005
aattattctt aatgtctgta a	aaaacgattt tcttctgtag	g aatgtttgac ttcgtattga	3065
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aaagtgcaaa aaaaaaaaa			3144

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	Ser					Ile					Met				gta Val 192	752
	Leu					Gln					Phe				ttg Leu 208	800
	Ile										Gln				cag Gln 224	848
	Gly														ggt Gly 240	896
cag Gln 241	Asn	att Ile	tat Tyr	aat Asn	gcc Ala 246	tgc Cys	tgt Cys	acc Thr	cta Leu	agg Arg 251	att Ile	gat Asp	ttt Phe	tcc Ser	aaa Lys 256	944
	Val				gta Val 262										tat Tyr 272	992
					cca Pro 278											1040
					ttt Phe 294											1088
gga Gly 305	gct Ala	ctg Leu	agt Ser	cct Pro	ttg Leu 310	gcc Ala	att Ile	cca Pro	aat Asn	gct Ala 315	gct Ala	gca Ala	gca Ala	gct Ala	gct Ala 320	1136
					cga Arg 326											1184
					gtt Val 342											1232
caa Gln 353	agt Ser	ctg Leu	ttt Phe	acc Thr	ctc Leu 358	ttc Phe	ggt Gly	gtt Val	tat Tyr	gga Gly 363	gat Asp	gtg Val	cag Gln	cgt Arg	gtg Val 368	1280
aag Lys 369	att Ile	tta Leu	tac Tyr	aat Asn	aag Lys 374	aaa · Lys	gac Asp	agc Ser	gct Ala	cta Leu 379	ata Ile	cag Gln	atg Met	gct Ala	gat Asp 384	1328
gga Gly 385	aac Asn	caa Gln	tca Ser	caa Gln	ctt Leu 390	gcc Ala	atg Met	aat Asn	cat His	ctt Leu 395	aat Asn	gga Gly	cag Gln	aaa Lys	atg Met 400	1376
tat	gga	aaa	att	att	cgt	gtt	act	ctg	tct	aaa	cat	cag	act	gta	cag	1424

Tyr Gly Lys Ile Ile Arg Val Thr Leu Ser Lys His Gln Thr Val Gln 401 406 411 416	
cta cct cga gag gga ctt gat gat caa ggg cta aca aaa gat ttt ggt Leu Pro Arg Glu Gly Leu Asp Asp Gln Gly Leu Thr Lys Asp Phe Gly 417 422 427 432	1472
aat tcc cca ttg cat cgt ttt aag aaa cct gga tcc aaa aat ttt caa Asn Ser Pro Leu His Arg Phe Lys Lys Pro Gly Ser Lys Asn Phe Gln 433 438 448	1520
aac att ttt cct cct tct gcc acc ctt cac cta tct aat atc cct cca Asn Ile Phe Pro Pro Ser Ala Thr Leu His Leu Ser Asn Ile Pro Pro 449 454 459 464	1568
tca gta gca gaa gag gat cta cga aca ctg ttc gct aac act ggg ggc Ser Val Ala Glu Glu Asp Leu Arg Thr Leu Phe Ala Asn Thr Gly Gly 465 470 475 480	1616
act gtg aaa gca ttt aag ttt ttt caa aga gat cac aaa atg gct ctt Thr Val Lys Ala Phe Lys Phe Phe Gln Arg Asp His Lys Met Ala Leu 481 486 491 496	1664
ctt cag atg gca aca gtg gaa gaa gct att cag gcc ttg att gat ctt Leu Gln Met Ala Thr Val Glu Glu Ala Ile Gln Ala Leu Ile Asp Leu 497 502 507 512	1712
cat aat tat aac ctt gga gaa aac cat cat ctg aga gtg tct ttc tcc His Asn Tyr Asn Leu Gly Glu Asn His His Leu Arg Val Ser Phe Ser 513 528	1760
aag tca aca att taa aaatgggaag atgaagattg ggggtgaatc acattgttca Lys Ser Thr Ile * 529	1815
atgtcatcac ctatttgact gttcagaaaa gtggggacca gagtttgatt ttttttgttt	1875
ttgttttttt ggggtttctt tttttttcc atgctgttat cattccttgg ttataaaatg	1935
aaatggcata tgtaaaggca gagttgttaa ctgctatatt tcatctgttc tatagggaag	1995
ccattttgtc tgtttaaaat ttcagtttaa ttttgctttt tttttttt ttttcctctt	2055
caacttagtt gacatacgtg ccttaaaaag gaaaactagt gttgctattg tgcatttacg	2115
ggaaaaaagg attggttgtt taggggcaca ctgttatatg gggtattaaa aatgtttaag	2175
gcaggggttt attaaaaaag gttaagtttt tttttaccac tgtttaggga ccttaatttt	2235
gaaatactgg gcttttcaac cttttttcca aattaaacca aaaagggtac caggatattg	2295
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176

tta	gac	ttg	agt	cac	aac	cat	ata	aaa	aag	ctt	cca	gct	aca	att	gga	630
Leu 182	Asp	Leu	Ser	His	Asn 187	His	Ile	Lys	Lys	Leu 192	Pro	Ala	Thr	Ile	Gly 197	
gac Asp 198	ctc Leu	ata Ile	cac His	ctt Leu	caa Gln 203	gaa Glu	ctt Leu	aac Asn	ctg Leu	aat Asn 208	gac Asp	aat Asn	cac His	ttg Leu	gag Glu 213	678
tca Ser 214	ttt Phe	agt Ser	gta Val	gcc Ala	ttg Leu 219	tgt Cys	cat His	tct Ser	aca Thr	ctc Leu 224	cag Gln	aag Lys	tca Ser	ctt Leu	cgg Arg 229	726
agt Ser 230	ttg Leu	gac Asp	ctc Leu	agc Ser	aag Lys 235	aac Asn	aaa Lys	atc Ile	aag Lys	gca Ala 240	ctc Leu	cct Pro	gtg Val	cag Gln	ttt Phe 245	774
tgc Cys 246	cag Gln	ctc Leu	cag Gln	gaa Glu	ctt Leu 251	aag Lys	aat Asn	tta Leu	aaa Lys	ctt Leu 256	gac Asp	gat Asp	aat Asn	gaa Glu	ttg Leu 261	822
att Ile 262	caa Gln	ttt Phe	cct Pro	tgc Cys	aag Lys 267	ata Ile	gga Gly	caa Gln	cta Leu	ata Ile 272	aac Asn	ctt Leu	cgc Arg	ttt Phe	ttg Leu 277	870
											agt Ser					918
											act Thr					966
											tta Leu					1014
											cca Pro					1062
											acc Thr					1110
gtt Val 358	tgt Cys	gga Gly	aga Arg	ttc Phe	tgt Cys 363	ctg Leu	aac Asn	tct Ser	ttc Phe	att Ile 368	caa Gln	gga Gly	act Thr	act Thr	acc Thr 373	1158
atg Met 374	aat Asn	ctg Leu	cat His	tct Ser	gtt Val 379	gcc Ala	cac His	act Thr	gtg Val	gtc Val 384	tta Leu	gta Val	gat Asp	aat Asn	ttg Leu 389	1206
ggt Gly 390	ggt Gly	act Thr	gaa Glu	gca Ala	cct Pro 395	att Ile	atc Ile	tct Ser	tat Tyr	ttc Phe 400	tgt Cys	tct Ser	cta Leu	ggc Gly	tgt Cys 405	1254
tat	gtt	aat	tcc	tct	gat	atg	tta	aag	taa	t gg	gtga	gaco	aga	aaaa	agaa	1305

Tyr Val 406	Asn Ser	Ser As		Leu	Lys	*								
atttcaa	taa caga	tcagtt	tgggg	tgcat	gta	tgat	ttt	gcag	gcgt	caa a	attg	gagtaa	1365	
gggaaga	ttt ctgt	atactt	gctgg	agagg	agg	gaato	gtgt	atag	gtta	ctc a	attt	agatga	1425	
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<2 <2	20> 21> CDS 22> (117		29)											
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gacccac	gcg tccg	gaggagg	ggggc	gacca	gca	agcca	aagc	ccag	gttc	cat (ctgc	cc	116	
atg gac Met Asp 1	atg gtg Met Val	g cac ct . His Le 5	g gtc eu Val	cat His	gcc Ala	aag Lys 10	aat Asn	gcc Ala	cag Gln	gct Ala	Leu 15	gcc Ala	164	
agc gac Ser Asp 17	cac gad His Asp	Tyr A	gg aca gg Thr	cag Gln	tat Tyr	cac His	aag Lys 27	ttc Phe	aca Thr	gca Ala	ctg Leu	ccc Pro 32	212	
gag gac Glu Asp 33	ctg aag Leu Lys	Met Al	cc tgg La Trp 88	gcc Ala	aag Lys	aaa Lys	gcc Ala 43	cat His	gcc Ala	ctg Leu	cag Gln	agt Ser 48	260	
gag ttg Glu Leu 49	cgc tac Arg Tyi	Lys Se	ca gac er Asp 54	ctg Leu	atc Ile	ggc Gly	atg Met 59	aag Lys	ggc Gly	ata Ile	gga Gly	tgg Trp 64	308	
ctg gcg Leu Ala 65	ctg aga Leu Arg	g Ser Pi	ca cag co Gln 70	atg Met	gag Glu	agt Ser	gca Ala 75	aag Lys	aag Lys	gct Ala	gga Gly	gaa Glu 80	356	
ctc atc Leu Ile 81	agc gag Ser Glu	ı Thr Ly	ag tac /s Tyr 86	cgt Arg	aaa Lys	aaa Lys	cca Pro 91	gac Asp	agt Ser	atc Ile	aag Lys	ttc Phe 96	404	
acc aca Thr Thr 97	gtg gtt Val Val	L Asp Se	cc cca er Pro 02	gac Asp	ctg Leu	gtt Val	cat His 107	gcc Ala	aag Lys	aac Asn	agc Ser	tat Tyr 112	452	

atg cac tgc aat gag cgc atg tat aga tct gga gat gca gaa tcc ctg

Met His Cys Asn Glu Arg Met Tyr Arg Ser Gly Asp Ala Glu Ser Leu

	Arg					Pro					. Phe				cgc Arg 144	548
	Asn										Arg				gag Glu 160	596
	Thr										Asp				ttc Phe 176	644
	Thr		cgg Arg													692
	Ala		ctg Leu													740
			cca Pro													788
			gag Glu													836
			aca Thr													884
			agt Ser													- 932
			gag Glu													980
Leu 289	Gln	Ser	gat Asp	Val	Lys 294	Tyr	Lys	Ser	Asp	Leu 299	Asn	Leu	Thr	Arg	Gly 304	1028
			acc Thr													1076
			ctg Leu													1124
cgg Arg 337	gly aaa	gca Ala	gaa Glu	gca Ala	gtg Val 342	gag Glu	gct Ala	gga Gly	gat Asp	cat Hi's 347	cag Gln	agt Ser	gly ggg	gag Glu	gtg Val 352	1172
aac	cca	gat	gcc	act	gag	att	ctg	cac	gtc	aaa	aag	aag	aag	gcc	ctg	1220

Asn Pro Asp Ala Thr Glu Ile Leu His Val Lys Lys Lys Lys Ala Leu 353 368	
ctg ttg tga gccatgt ccaccctgat tcctgagagg ccagagagga agtttgttca Leu Leu * 369	1276
ccagagacag gcttcagatg gctttgattt cggcaagctg gaatggccca ccagcccatc	1336
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gacaagetge ggeteageee caetgacett ggeteetgee egeeetgegg eeeetgeeee	180
atcccgaagc cggcagccag aggcaggcgc cagagtcaag actggggcaa gagtgacgag	240
aggetgetae aageegtgga aaacaaegat geaeeteggg tggeegeeet categeeege	300
aaggggctgg tgcccacgaa gctagacccc gagggcaagt ccgcgttcca cctggcggcc	360
atgeggggtg eggeeagetg tetggaggtg atgatagete atggeageaa tgteatgage	420
geggaegggg caggttacaa tgeeeteeae etggeegeea aataegggea eecacagtge	480
ttgaagcaac tactgcaggc ttcctgcgtg gtggacgtcg tggacagcag cgggtggact	540
gccctacacc atg cag ggg tgg att gcc cta gca gcg gct ggt ggc tgt Met Gln Gly Trp Ile Ala Leu Ala Ala Ala Gly Gly Cys 1 5 10	589
ctc tcc tgc tca gag gtg ctc tgc tcc ttt aag gca cat cta aac ccc Leu Ser Cys Ser Glu Val Leu Cys Ser Phe Lys Ala His Leu Asn Pro 14 19 24 29	637
caa gat cgg tca ggc gca aca ccc ctc att ata gca gct cag atg tgt Gln Asp Arg Ser Gly Ala Thr Pro Leu Ile Ile Ala Ala Gln Met Cys 30 40 45	685
cac aca gac ctg tgc cgt ctc cta ctg cag caa ggg gct gcc gcg aac His Thr Asp Leu Cys Arg Leu Leu Cln Gln Gly Ala Ala Ala Asn 46 51 . 56 61	733

gat Asp 62	cag Gln	gac Asp	ctg Leu	caa Gln	ggc Gly 67	Arg	acg Thr	gcc Ala	ctg Leu	atg Met 72	ctg Leu	gcc Ala	tgt Cys	gag Glu	999 Gly 77	781
gcc Ala 78	agc Ser	ccc Pro	gaa Glu :	aca Thr	gtg Val 83	gag Glu	gtc Val	ctg Leu	ctg Leu	cag Gln 88	ggc Gly	gga Gly	gcc Ala	cag Gln	ccg Pro 93	829
ggc Gly 94	atc Ile	acc Thr	gat Asp	gcg Ala	ctg Leu 99	Gly aaa	cag Gln	gac Asp	gcg Ala	gct Ala 104	cac His	tat Tyr	ggc Gly	gcc Ala	ctg Leu 109	877
gcg Ala 110	Gly aaa	gac Asp	aaa Lys	ctc Leu	atc Ile 115	ctg Leu	cac His	ctt Leu	ctg Leu	caa Gln 120	gag [†] Glu	gcg Ala	gcc Ala	cag Gln	cgc Arg 125	925
ccc Pro 126	tcc Ser	cca Pro	ccc Pro	agc Ser	gcc Ala 131	ctc Leu	aca Thr	gag Glu	gat Asp	gat Asp 136	tca Ser	ggc Gly	gag Glu	gcg Ala	tca Ser 141	973
tct Ser 142	cag Gln	aac Asn	tct Ser	atg Met	tcc Ser 147	agc Ser	cat His	gga Gly	aag Lys	cag Gln 152	ggg Gly	gcc Ala	ccc Pro	aag Lys	aag Lys 157	1021
cgg Arg 158	aag Lys	gcg Ala	cct Pro	cca Pro	cct Pro 163	ccc Pro	gcc Ala	agc Ser	att Ile	ccc Pro 168	atg Met	ccg Pro	gat Asp	gat Asp	cga Arg 173	1069
gat Asp 174	gcc Ala	tat Tyr	gag Glu	gag Glu	atc Ile 179	gtg Val	agg Arg	ctg Leu	cgg Arg	cag Gln 184	gag Glu	agg Arg	ggc Gly	cgc Arg	ctc Leu 189	1117
ctg Leu 190	cag Gln	aag Lys	atc Ile	cgg Arg	ggc Gly 195	ctg Leu	gaa Glu	cag Gln	cac His	aag Lys 200	gaa Glu	cgg Arg	agg Arg	cag Gln	cag Gln 205	1165
gag Glu 206	tcc Ser	ccg Pro	gag Glu	gcc Ala	agc Ser 211	tcc Ser	ctg Leu	cac His	atc Ile	ctg Leu 216	gag Glu	aga Arg	cag Gln	gtg Val	caa Gln 221	1213
gag Glu 222	cta Leu	cag Gln	cag Gln	ttg Leu	ctg Leu 227	gtg Val	gag Glu	aga Arg	caa Gln	gag Glu 232	gag Glu	aag Lys	gag Glu	agc Ser	ctg Leu 237	1261
gga Gly 238	cgg Arg	gag Glu	gtg Val	gag Glu	agt Ser 243	ttg Leu	cag Gln	agc Ser	cgg Arg	ctg Leu 248	tcc Ser	ctg Leu	ctg Leu	gag Glu	aac Asn 253	1309
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ggt Gly 270	gag Glu	ctg Leu	cct Pro	gac Asp	ctt Leu 275	cca Pro	Gly 999	gcc Ala	gag Glu	gtg Val 280	ctg Leu	ctg Leu	tcc Ser	aga Arg	caa Gln 285	1405

ctc agt ccg tcg gcc cag gaa cac ctg gcc tcg ctg cag gaa cag gtg Leu Ser Pro Ser Ala Gln Glu His Leu Ala Ser Leu Gln Glu Gln Val 286 291 296 301	1453
gct gtg ctc acc aga cag aac cag gaa ctg atg gag aag gtc cag atc Ala Val Leu Thr Arg Gln Asn Gln Glu Leu Met Glu Lys Val Gln Ile 302 307 312 317	1501
ctg gag aac ttt gag aag gac gag aca cag atg gaa gtg gaa gct ttg Leu Glu Asn Phe Glu Lys Asp Glu Thr Gln Met Glu Val Glu Ala Leu 318 323 328 333	1549
gca gag gtc att cct ctt gcc ctt atg aat ctc tct cgg gcc gag ttt Ala Glu Val Ile Pro Leu Ala Leu Met Asn Leu Ser ¹³ Arg Ala Glu Phe 334 339 344 349	1597
gac cat ttc gca ggc aca cgc ttg agc ctt gag gcc ctg agg cag cag Asp His Phe Ala Gly Thr Arg Leu Ser Leu Glu Ala Leu Arg Gln Gln 350 355 360 365	1645
gag aca cga gat gtc ccc aga gaa tag gggcc acctgtggga gagtgaggtg Glu Thr Arg Asp Val Pro Arg Glu * 366 371	1697
ctgagccacg ccact .	1712
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ttaaaaatgg tgggtattcg atttttctct cgtttttgtc aattcgattg tccacagtta	180
tggtcacgcc cgtcaccaac ggcttaccgg atcgtttcac agtaagccct acgatgaaat	240
gtgacaaaaa ttgactttat tcagcaaaaa tgaaaatcag cccgctgaat atgtaacatt	300
aattaacccg aacaatacgg gcaaaaaaca ttcatgataa aaatatttat tggtcattat	360
attaacgtat tttatagcac tgccgatatc acgctcaaaa aacaaccact gctatttta	420
gcaaagctta tggtatactc cgccgcctta acatttttca ccgcaaattt/tcattgcaac	480
atgacgagga aaataa atg agt acg cct gat ttt tct act gcc gag aat Met Ser Thr Pro Asp Phe Ser Thr Ala Glu Asn	529

													atg Met			577
													gtg Val			625
_	_	_		_			_		_	_	_		cag Gln	-	_	673
													cgt Arg		taa * 75	721
tata	aaac	cg g	gctga	atago	g to	geett	tcag	g ccg	gttt	ttg	cato	tggd	cac c	gcago	gacaga	781
aaag	jagat	ca a	atta	agco	a gt	agco	gccg	g ct								813

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agcaattttc aatgcggcaa ccggcaaggc tgaccgtgta ggctttagat tcgaagacgg

taaaaaagtc cgtttcttca agtctaacag cgaaactatc aagtaatttg gagtagtacg

atg gcg aaa ctg cat gat tac tac aaa gac gaa gta gtt aaa aaa ctc Met Ala Lys Leu His Asp Tyr Tyr Lys Asp Glu Val Val Lys Lys Leu 480

															gag Glu 32	636
					atg Met 38										aaa Lys 48	684
					gca Ala 54										aaa Lys 64	732
_	_				gca Ala 70	_			_	_					_	780
					ggc Gly 86											828
					cgc Arg 102											876
					tcc Ser 118											924
					gag Glu 134											972
					cgt Arg 150											1020
					ggc Gly 166											1068
	cgc Arg		taa *	ggt	aggg	ıttac	ta a	atgg	ıctaa	ig ca	atca	ıatga	aaç	cacc	ıcga	1123
agta	aaac	gc g	ıtago	ttta	g ct	gata	aata	ctt	cgcg	jaaa	cgcg	ıctga	ac t	gaaa	ıgcgat	1183
cato	tato	gat g	ıtgaa	cgct	t cc	gacg	jaa									1211

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<211> 1327

<212> DNA

<213> Homo sapiens

<220>

<221> CDS <222> (251)..(1300)

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Lys 174	Ser	Glu	Glu	Ile	Gly 179	Lys	Lys	Gln	Leu	Ser 184	Lys	Asp	Ser	Ile	Leu 189	•
	ctg Leu															865
	gct Ala															913
	gtt Val															961
gta Val 238	ggc Gly	atg Met	gtt Val	gct Ala	cag Gln 243	cca Pro	gga Gly	gct Ala	tct Ser	999 Gly 248	atg Met	gtt Val	gcc Ala	ccc Pro	atg Met 253	1009
	atg Met															1057
	ccg Pro															1105
	gca Ala															1153
	caa Gln															1201
	tat Tyr															1249
	aat Asn															1297
taa * 350	aaac	aaaa	ca c	ctgt	aaaa	aa aa	aaaa	ıa								1327